SEQUENCE LISTING

| <110> | CAN | FIELD, Will | iam | | • | | | | | | | | |
|---------|-----------------------------------|-------------|------------|------------|------------|------------|------|--|--|--|--|--|--|
| <120> | SOLUBLE GLCNAC PHOSPHOTRANSFERASE | | | | | | | | | | | | |
| <130> | 203515US77 | | | | | | | | | | | | |
| <160> | 38 | | | | | | | | | | | | |
| <170> | PatentIn version 3.1 | | | | | | | | | | | | |
| | 1 3600 DNA hybi | | | | | | | | | | | | |
| <400> | 1 | | • | | | | | | | | | | |
| atggaga | cag | acacactcct | gctatgggta | ctgctgctct | gggttccagg | ttccactggt | 60 | | | | | | |
| gacgaag | atc | aggtagatcc | gcggttaatc | gacggtaagc | ttagccgaga | tcaataccat | 120 | | | | | | |
| gttttgt | ttg | attcctatag | agacaatatt | gctggaaagt | cctttcagaa | tcggctttgt | 180 | | | | | | |
| ctgccca | tgc | cgattgacgt | tgtttacacc | tgggtgaatg | gcacagatct | tgaactactg | 240 | | | | | | |
| aaggaac | tac | agcaggtcag | agaacagatg | gaggaggagc | agaaagcaat | gagagaaatc | 300 | | | | | | |
| cttggga | aaa | acacaacgga | acctactaag | aagagtgaga | agcagttaga | gtgtttgcta | 360 | | | | | | |
| acacact | gca | ttaaggtgcc | aatgcttgtc | ctggacccag | ccctgccagc | caacatcacc | 420 | | | | | | |
| ctgaagg | acc | tgccatctct | ttatccttct | tttcattctg | ccagtgacat | tttcaatgtt | 480 | | | | | | |
| gcaaaac | caa | aaaacccttc | taccaatgtc | tcagttgttg | tttttgacag | tactaaggat | 540 | | | | | | |
| gttgaag | atg | cccactctgg | actgcttaaa | ggaaatagca | gacagacagt | atggagggc | 600 | | | | | | |
| tacttga | caa | cagataaaga | agtccctgga | ttagtgctaa | tgcaagattt | ggctttcctg | 660 | | | | | | |
| agtggat | ttc | caccaacatt | caaggaaaca | aatcaactaa | aaacaaaatt | gccagaaaat | 720 | | | | | | |
| ctttcct | cta | aagtcaaact | gttgcagttg | tattcagagg | ccagtgtagc | gcttctaaaa | 780 | | | | | | |
| ctgaata | acc | ccaaggattt | tcaagaattg | aataagcaaa | ctaagaagaa | catgaccatt | 840 | | | | | | |
| gatggaa | aag | aactgaccat | aagtcctgca | tatttattat | gggatctgag | cgccatcagc | 900 | | | | | | |
| cagtcta | agc | aggatgaaga | catctctgcc | agtcgttttg | aagataacga | agaactgagg | 960 | | | | | | |
| tactcat | tgc | gatctatcga | gaggcatgca | ccatgggttc | ggaatatttt | cattgtcacc | 1020 | | | | | | |
| aacgggc | aga | ttccatcctg | gctgaacctt | gacaatcctc | gagtgacaat | agtaacacac | 1080 | | | | | | |
| caggatg | ttt | ttcgaaattt | gagccacttg | cctaccttta | gttcacctgc | tattgaaagt | 1140 | | | | | | |
| cacgttc | atc | gcatcgaagg | gctgtcccag | aagtttattt | acctaaatga | tgatgtcatg | 1200 | | | | | | |

| tttgggaagg atgtctggcc | agatgatttt | tacagtcact | ccaaaggcca | gaaggtttat | 1260 |
|-----------------------|------------|------------|------------|------------|------|
| ttgacatggc ctgtgccaaa | ctgtgccgag | ggctgcccag | gttcctggat | taaggatggc | 1320 |
| tattgtgaca aggettgtaa | taattcagcc | tgcgattggg | atggtgggga | ttgctctgga | 1380 |
| aacagtggag ggagtcgcta | tattgcagga | ggtggaggta | ctgggagtat | tggagttgga | 1440 |
| cagccctggc agtttggtgg | aggaataaac | agtgtctctt | actgtaatca | gggatgtgcg | 1500 |
| aatteetgge tegetgataa | gttctgtgac | caagcatgca | atgtcttgtc | ctgtgggttt | 1560 |
| gatgctggcg actgtgggca | agatcatttt | catgaattgt | ataaagtgat | ccttctccca | 1620 |
| aaccagactc actatattat | tccaaaaggt | gaatgcctgc | cttatttcag | ctttgcagaa | 1680 |
| gtagccaaaa gaggagttga | aggtgcctat | agtgacaatc | caataattcg | acatgcttct | 1740 |
| attgccaaca agtggaaaac | catccacctc | ataatgcaca | gtggaatgaa | tgccaccaca | 1800 |
| atacatttta atctcacgtt | tcaaaataca | aacgatgaag | agttcaaaat | gcagataaca | 1860 |
| gtggaggtgg acacaaggga | gggaccaaaa | ctgaattcta | cggcccagaa | gggttacgaa | 1920 |
| aatttagtta gtcccataac | acttcttcca | gaggcggaaa | tcctttttga | ggatattccc | 1980 |
| aaagaaaaac gcttcccgaa | gtttaagaga | catgatgtta | actcaacaag | gagagcccag | 2040 |
| gaagaggtga aaattcccct | ggtaaatatt | tcactccttc | caaaagacgc | ccagttgagt | 2100 |
| ctcaatacct tggatttgca | actggaacat | ggagacatca | ctttgaaagg | atacaatttg | 2160 |
| tccaagtcag ccttgctgag | atcatttctg | atgaactcac | agcatgctaa | aataaaaaat | 2220 |
| caagctataa taacagatga | aacaaatgac | agtttggtgg | ctccacagga | aaaacaggtt | 2280 |
| cataaaagca tcttgccaaa | cagcttagga | gtgtctgaaa | gattgcagag | gttgactttt | 2340 |
| cctgcagtga gtgtaaaagt | gaatggtcat | gaccagggtc | agaatccacc | cctggacttg | 2400 |
| gagaccacag caagatttag | agtggaaact | cacacccaaa | aaaccatagg | cggaaatgtg | 2460 |
| acaaaagaaa agcccccatc | tctgattgtt | ccactggaaa | gccagatgac | aaaagaaaag | 2520 |
| aaaatcacag ggaaagaaaa | agagaacagt | agaatggagg | aaaatgctga | aaatcacata | 2580 |
| ggcgttactg aagtgttact | tggaagaaag | ctgcagcatt | acacagatag | ttacttgggc | 2640 |
| tttttgccat gggagaaaaa | aaagtatttc | ctagatette | tcgacgaaga | agagtcattg | 2700 |
| aagacacaat tggcctactt | cactgatagc | aagaatagag | ccagatacaa | gagagataca | 2760 |
| tttgcagatt ccctcagata | tgtaaataaa | attctaaata | gcaagtttgg | attcacatcg | 2820 |
| cggaaagtcc ctgctcacat | gcctcacatg | attgaccgga | ttgttatgca | agaactgcaa | 2880 |

| gatatgttcc | ctgaagaatt | tgacaagacg | tcatttcaca | aagtgcgcca | ttctgaggat | 2940 |
|------------|------------|------------|------------|------------|------------|------|
| atgcagtttg | ccttctctta | tttttattat | ctcatgagtg | cagtgcagcc | actgaatata | 3000 |
| tctcaagtct | ttgatgaagt | tgatacagat | caatctggtg | tcttgtctga | cagagaaatc | 3060 |
| cgaacactgg | ctaccagaat | tcacgaactg | ccgttaagtt | tgcaggattt | gacaggtctg | 3120 |
| gaacacatgc | taataaattg | ctcaaaaatg | cttcctgctg | atatcacgca | gctaaataat | 3180 |
| attccaccaa | ctcaggaatc | ctactatgat | cccaacctgc | caccggtcac | taaaagtcta | 3240 |
| gtaacaaact | gtaaaccagt | aactgacaaa | atccacaaag | catataagga | caaaaacaaa | 3300 |
| tataggtttg | aaatcatggg | agaagaagaa | atcgctttta | aaatgattcg | taccaacgtt | 3360 |
| tctcatgtgg | ttggccagtt | ggatgacata | agaaaaaacc | ctaggaagtt | tgtttgcctg | 3420 |
| aatgacaaca | ttgaccacaa | tcataaagat | gctcagacag | tgaaggctgt | tctcagggac | 3480 |
| ttctatgaat | ccatgttccc | cataccttcc | caatttgaac | tgccaagaga | gtatcgaaac | 3540 |
| cgtttccttc | atatgcatga | gctgcaggaa | tggagggctt | atcgagacaa | attgaagtag | 3600 |

<210> 2

<211> 1199

<212> PRT

<213> hybrid

<400> 2

Met Glu Thr Asp Thr Leu Leu Leu Trp Val Leu Leu Leu Trp Val Pro 1 5 10 15

Gly Ser Thr Gly Asp Glu Asp Gln Val Asp Pro Arg Leu Ile Asp Gly 20 25 30

Lys Leu Ser Arg Asp Gln Tyr His Val Leu Phe Asp Ser Tyr Arg Asp 35 40 45

Asn Ile Ala Gly Lys Ser Phe Gln Asn Arg Leu Cys Leu Pro Met Pro 50 55 60

Ile Asp Val Val Tyr Thr Trp Val Asn Gly Thr Asp Leu Glu Leu Leu 65 70 75 80

Lys Glu Leu Gln Gln Val Arg Glu Gln Met Glu Glu Glu Gln Lys Ala 85 90 95

Met Arg Glu Ile Leu Gly Lys Asn Thr Thr Glu Pro Thr Lys Lys Ser

100 105 110

Glu Lys Gln Leu Glu Cys Leu Leu Thr His Cys Ile Lys Val Pro Met 115 120 125

Leu Val Leu Asp Pro Ala Leu Pro Ala Asn Ile Thr Leu Lys Asp Leu 130 135 140

Pro Ser Leu Tyr Pro Ser Phe His Ser Ala Ser Asp Ile Phe Asn Val 145 150 155 160

Ala Lys Pro Lys Asn Pro Ser Thr Asn Val Ser Val Val Val Phe Asp 165 170 175

Ser Thr Lys Asp Val Glu Asp Ala His Ser Gly Leu Leu Lys Gly Asn 180 185 190

Ser Arg Gln Thr Val Trp Arg Gly Tyr Leu Thr Thr Asp Lys Glu Val

Pro Gly Leu Val Leu Met Gln Asp Leu Ala Phe Leu Ser Gly Phe Pro 210 215 220

Pro Thr Phe Lys Glu Thr Asn Gln Leu Lys Thr Lys Leu Pro Glu Asn 225 230 235 240

Leu Ser Ser Lys Val Lys Leu Gln Leu Tyr Ser Glu Ala Ser Val 245 250 255

Ala Leu Leu Lys Leu Asn Asn Pro Lys Asp Phe Gln Glu Leu Asn Lys 260 265 270

Gln Thr Lys Lys Asn Met Thr Ile Asp Gly Lys Glu Leu Thr Ile Ser 275 280 285

Pro Ala Tyr Leu Leu Trp Asp Leu Ser Ala Ile Ser Gln Ser Lys Gln 290 295 300

Asp Glu Asp Ile Ser Ala Ser Arg Phe Glu Asp Asn Glu Glu Leu Arg 305 310 315 320

Tyr Ser Leu Arg Ser Ile Glu Arg His Ala Pro Trp Val Arg Asn Ile 325 330 335 Phe Ile Val Thr Asn Gly Gln Ile Pro Ser Trp Leu Asn Leu Asp Asn 340 345 350

Pro Arg Val Thr Ile Val Thr His Gln Asp Val Phe Arg Asn Leu Ser 355 360 365

His Leu Pro Thr Phe Ser Ser Pro Ala Ile Glu Ser His Val His Arg 370 375 380

Ile Glu Gly Leu Ser Gln Lys Phe Ile Tyr Leu Asn Asp Asp Val Met 385 390 395 400

Phe Gly Lys Asp Val Trp Pro Asp Asp Phe Tyr Ser His Ser Lys Gly 405 410 415

Gln Lys Val Tyr Leu Thr Trp Pro Val Pro Asn Cys Ala Glu Gly Cys 420 425 430

Pro Gly Ser Trp Ile Lys Asp Gly Tyr Cys Asp Lys Ala Cys Asn Asn 435 440 445

Ser Ala Cys Asp Trp Asp Gly Gly Asp Cys Ser Gly Asn Ser Gly Gly 450 455 460

Ser Arg Tyr Ile Ala Gly Gly Gly Gly Thr Gly Ser Ile Gly Val Gly 465 470 475 480

Gln Pro Trp Gln Phe Gly Gly Gly Ile Asn Ser Val Ser Tyr Cys Asn 485 490 495

Gln Gly Cys Ala Asn Ser Trp Leu Ala Asp Lys Phe Cys Asp Gln Ala 500 505 510

Cys Asn Val Leu Ser Cys Gly Phe Asp Ala Gly Asp Cys Gly Gln Asp 515 520 525

His Phe His Glu Leu Tyr Lys Val Ile Leu Leu Pro Asn Gln Thr His 530 540

Tyr Ile Ile Pro Lys Gly Glu Cys Leu Pro Tyr Phe Ser Phe Ala Glu 545 550 555 560 Val Ala Lys Arg Gly Val Glu Gly Ala Tyr Ser Asp Asn Pro Ile Ile 565 570 575

Arg His Ala Ser Ile Ala Asn Lys Trp Lys Thr Ile His Leu Ile Met 580 585 590

His Ser Gly Met Asn Ala Thr Thr Ile His Phe Asn Leu Thr Phe Gln 595 600 605

Asn Thr Asn Asp Glu Glu Phe Lys Met Gln Ile Thr Val Glu Val Asp 610 615 620

Thr Arg Glu Gly Pro Lys Leu Asn Ser Thr Ala Gln Lys Gly Tyr Glu 625 630 635 640

Asn Leu Val Ser Pro Ile Thr Leu Leu Pro Glu Ala Glu Ile Leu Phe 645 650 655

Glu Asp Ile Pro Lys Glu Lys Arg Phe Pro Lys Phe Lys Arg His Asp 660 665 670

Val Asn Ser Thr Arg Arg Ala Glu Glu Val Lys Ile Pro Leu Val 675 680 685

Asn Ile Ser Leu Leu Pro Lys Asp Ala Gln Leu Ser Leu Asn Thr Leu 690 695 700

Asp Leu Gln Leu Glu His Gly Asp Ile Thr Leu Lys Gly Tyr Asn Leu 705 710 715 720

Ser Lys Ser Ala Leu Leu Arg Ser Phe Leu Met Asn Ser Gln His Ala 725 730 735

Lys Ile Lys Asn Gln Ala Ile Ile Thr Asp Glu Thr Asn Asp Ser Leu 740 745 750

Val Ala Pro Gln Glu Lys Gln Val His Lys Ser Ile Leu Pro Asn Ser 755 760 765

Leu Gly Val Ser Glu Arg Leu Gln Arg Leu Thr Phe Pro Ala Val Ser 770 775 780

odezasa

<u></u>

Val Lys Val Asn Gly His Asp Gln Gly Gln Asn Pro Pro Leu Asp Leu 790

Glu Thr Thr Ala Arg Phe Arg Val Glu Thr His Thr Gln Lys Thr Ile 810 805

Gly Gly Asn Val Thr Lys Glu Lys Pro Pro Ser Leu Ile Val Pro Leu 825 820

Glu Ser Gln Met Thr Lys Glu Lys Lys Ile Thr Gly Lys Glu Lys Glu 845 840 835

Asn Ser Arg Met Glu Glu Asn Ala Glu Asn His Ile Gly Val Thr Glu 850

Val Leu Leu Gly Arg Lys Leu Gln His Tyr Thr Asp Ser Tyr Leu Gly 870

Phe Leu Pro Trp Glu Lys Lys Lys Tyr Phe Leu Asp Leu Leu Asp Glu 885

Glu Glu Ser Leu Lys Thr Gln Leu Ala Tyr Phe Thr Asp Ser Lys Asn 910 905 900

Arg Ala Arg Tyr Lys Arg Asp Thr Phe Ala Asp Ser Leu Arg Tyr Val 920

Asn Lys Ile Leu Asn Ser Lys Phe Gly Phe Thr Ser Arg Lys Val Pro 935

Ala His Met Pro His Met Ile Asp Arg Ile Val Met Gln Glu Leu Gln 950

Asp Met Phe Pro Glu Glu Phe Asp Lys Thr Ser Phe His Lys Val Arg 970

His Ser Glu Asp Met Gln Phe Ala Phe Ser Tyr Phe Tyr Tyr Leu Met 990 980

Ser Ala Val Gln Pro Leu Asn Ile Ser Gln Val Phe Asp Glu Val Asp 1000

Thr Asp Gln Ser Gly Val Leu Ser Asp Arg Glu Ile Arg Thr Leu

1010 1015 1020

Ala Thr Arg Ile His Glu Leu Pro Leu Ser Leu Gln Asp Leu Thr 1025 1030 1035

Gly Leu Glu His Met Leu Ile Asn Cys Ser Lys Met Leu Pro Ala 1040 1045 1050

Asp Ile Thr Gln Leu Asn Asn Ile Pro Pro Thr Gln Glu Ser Tyr 1055 1060 1065

Tyr Asp Pro Asn Leu Pro Pro Val Thr Lys Ser Leu Val Thr Asn 1070 1075 1080

Cys Lys Pro Val Thr Asp Lys Ile His Lys Ala Tyr Lys Asp Lys 1085 1090 1095

Asn Lys Tyr Arg Phe Glu Ile Met Gly Glu Glu Glu Ile Ala Phe 1100 1105 1110

Lys Met Ile Arg Thr Asn Val Ser His Val Val Gly Gln Leu Asp 1115 1120 1125

Asp Ile Arg Lys Asn Pro Arg Lys Phe Val Cys Leu Asn Asp Asn 1130 1135 1140

Ile Asp His Asn His Lys Asp Ala Gln Thr Val Lys Ala Val Leu 1145 1150 1155

Arg Asp Phe Tyr Glu Ser Met Phe Pro Ile Pro Ser Gln Phe Glu 1160 1165 1170

Leu Pro Arg Glu Tyr Arg Asn Arg Phe Leu His Met His Glu Leu 1175 1180 1185

Gln Glu Trp Arg Ala Tyr Arg Asp Lys Leu Lys 1190 1195

<210> 3

<211> 5597

<212> DNA

<213> Homo sapiens

<400> 3

cggagccgag cgggcgtccg tcgccggagc tgcaatgagc ggcgcccgga ggctgtgacc 60 tgegegegge ggeeegaeeg gggeeeetga atggeggete getgaggegg eggeggegge 120 ggcggctcag gctcctcggg gcgtggcgtg gcggtgaagg ggtgatgctg ttcaagctcc 180 tgcagagaca aacctatacc tgcctgtccc acaggtatgg gctctacgtg tgcttcttgg 240 gegtegttgt caccategte teegeettee agtteggaga ggtggttetg gaatggagee 300 360 gagatcaata ccatgttttg tttgattcct atagagacaa tattgctgga aagtcctttc agaatcggct ttgtctgccc atgccgattg acgttgttta cacctgggtg aatggcacag 420 atcttgaact actgaaggaa ctacagcagg tcagagaaca gatggaggag gagcagaaag 480 caatgagaga aatccttggg aaaaacacaa cggaacctac taagaagagt gagaagcagt 540 tagagtgttt gctaacacac tgcattaagg tgccaatgct tgtactggac ccagccctgc 600 cagccaacat caccctgaag gacgtgccat ctctttatcc ttctttcat tctgccagtg 660 acattttcaa tgttgcaaaa ccaaaaaacc cttctaccaa tgtctcagtt gttgtttttg 720 acagtactaa ggatgttgaa gatgcccact ctggactgct taaaggaaat agcagacaga 780 cagtatggag ggggtacttg acaacagata aagaagtccc tggattagtg ctaatgcaag 840 atttggcttt cctgagtgga tttccaccaa cattcaagga aacaaatcaa ctaaaaacaa 900 aattgccaga aaatctttcc tctaaagtca aactgttgca gttgtattca gaggccagtg 960 tagcgcttct aaaactgaat aaccccaagg attttcaaga attgaataag caaactaaga 1020 agaacatgac cattgatgga aaagaactga ccataagtcc tgcatattta ttatgggatc 1080 tgagcgccat cagccagtct aagcaggatg aagacatctc tgccagtcgt tttgaagata 1140 acgaagaact gaggtactca ttgcgatcta tcgagaggca tgcaccatgg gttcggaata 1200 ttttcattgt caccaacggg cagattccat cctggctgaa ccttgacaat cctcgagtga 1260 caatagtaac acaccaggat gtttttcgaa atttgagcca cttgcctacc tttagttcac 1320 1380 ctgctattga aagtcacatt catcgcatcg aagggctgtc ccagaagttt atttacctaa atgatgatgt catgtttggg aaggatgtct ggccagatga tttttacagt cactccaaag 1440 gccagaaggt ttatttgaca tggcctgtgc caaactgtgc cgagggctgc ccaggttcct 1500 ggattaagga tggctattgt gacaaggctt gtaataattc agcctgcgat tgggatggtg 1560 gggattgctc tggaaacagt ggagggagtc gctatattgc aggaggtgga ggtactggga 1620 gtattggagt tggacacccc tggcagtttg gtggaggaat aaacagtgtc tcttactgta 1680 atcagggatg tgcgaattcc tggctcgctg ataagttctg tgaccaagca tgcaatgtct 1740 tgtcctgtgg gtttgatgct ggcgactgtg ggcaagatca ttttcatgaa ttgtataaag 1800 tgatccttct cccaaaccag actcactata ttattccaaa aggtgaatgc ctgccttatt 1860 tcagctttgc agaagtagcc aaaagaggag ttgaaggtgc ctatagtgac aatccaataa 1920 ttcgacatgc ttctattgcc aacaagtgga aaaccatcca cctcataatg cacagtggaa 1980 tgaatgccac cacaatacat tttaatctca cgtttcaaaa tacaaacgat gaagagttca 2040 aaatgcagat aacagtggag gtggacacaa gggagggacc aaaactgaat tctacggccc 2100 agaagggtta cgaaaattta gttagtccca taacacttct tccagaggcg gaaatccttt 2160 ttgaggatat tcccaaagaa aaacgcttcc cgaagtttaa gagacatgat gttaactcaa 2220 caaggagagc ccaggaagag gtgaaaattc ccctggtaaa tatttcactc cttccaaaag 2280 acgcccagtt gagtctcaat accttggatt tgcaactgga acatggagac atcactttga 2340 aaggatacaa tttgtccaag tcagccttgc tgagatcatt tctgatgaac tcacagcatg 2400 2460 ctaaaataaa aaatcaagct ataataacag atgaaacaaa tgacagtttg gtggctccac aggaaaaaca ggttcataaa agcatcttgc caaacagctt aggagtgtct gaaagattgc 2520 2580 agaggttgac ttttcctgca gtgagtgtaa aagtgaatgg tcatgaccag ggtcagaatc caccctgga cttggagacc acagcaagat ttagagtgga aactcacacc caaaaaacca 2640 2700 taggcggaaa tgtgacaaaa gaaaagcccc catctctgat tgttccactg gaaagccaga tgacaaaaga aaagaaaatc acagggaaag aaaaagagaa cagtagaatg gaggaaaatg 2760 ctgaaaatca cataggcgtt actgaagtgt tacttggaag aaagctgcag cattacacag 2820 atagttactt gggctttttg ccatgggaga aaaaaaagta tttccaagat cttctcgacg 2880 aagaagagtc attgaagaca caattggcat acttcactga tagcaaaaat actgggaggc 2940 aactaaaaga tacatttgca gattccctca gatatgtaaa taaaattcta aatagcaagt 3000 3060 ttggattcac atcgcggaaa gtccctgctc acatgcctca catgattgac cggattgtta tgcaagaact gcaagatatg ttccctgaag aatttgacaa gacgtcattt cacaaagtgc 3120 3180 agccactgaa tatatctcaa gtctttgatg aagttgatac agatcaatct ggtgtcttgt 3240 ctgacagaga aatccgaaca ctggctacca gaattcacga actgccgtta agtttgcagg 3300 3360 atttgacagg tctggaacac atgctaataa attgctcaaa aatgcttcct gctgatatca cgcagctaaa taatattcca ccaactcagg aatcctacta tgatcccaac ctgccaccgg 3420



| tcagccttct | gagtagctgg | gactacaggt | gcatgccacc | acaccctgct | aatttttgta | 5220 |
|------------|------------|------------|------------|------------|------------|------|
| tttttagtag | agacgggggt | tccaccatat | tggtcaggct | tatcttgaac | tcctgacctc | 5280 |
| aggtgatcca | cctgcctctg | cctcccaaag | tgctgagatt | acaggcataa | gccagtgcac | 5340 |
| ccagccgaga | attagtattt | ttatgtatgg | ttaaaccttg | gcgtctagcc | atattttatg | 5400 |
| tcataataca | atggatttgt | gaagagcaga | ttccatgagt | aactctgaca | ggtattttag | 5460 |
| atcatgatct | caacaatatt | cctcccaaat | ggcatacatc | ttttgtacaa | agaacttgaa | 5520 |
| atgtaaatac | tgtgtttgtg | ctgtaagagt | tgtgtatttc | aaaaactgaa | atctcataaa | 5580 |
| aagttaaatt | ttgaaaa | | | | | 5597 |

<210> 4 <211> 928 <212> PRT <213> Homo sapiens

<400> 4

Met Leu Phe Lys Leu Leu Gln Arg Gln Thr Tyr Thr Cys Leu Ser His 1 5 10 15

Arg Tyr Gly Leu Tyr Val Cys Phe Leu Gly Val Val Val Thr Ile Val 20 25 30

Ser Ala Phe Gln Phe Gly Glu Val Val Leu Glu Trp Ser Arg Asp Gln 35 40 45

Tyr His Val Leu Phe Asp Ser Tyr Arg Asp Asn Ile Ala Gly Lys Ser 50 55 60

Phe Gln Asn Arg Leu Cys Leu Pro Met Pro Ile Asp Val Val Tyr Thr 65 70 75 80

Trp Val Asn Gly Thr Asp Leu Glu Leu Leu Lys Glu Leu Gln Gln Val 85 90 95

Arg Glu Gln Met Glu Glu Glu Gln Lys Ala Met Arg Glu Ile Leu Gly 100 105 110

Lys Asn Thr Thr Glu Pro Thr Lys Lys Ser Glu Lys Gln Leu Glu Cys 115 120 125 Leu Leu Thr His Cys Ile Lys Val Pro Met Leu Val Leu Asp Pro Ala 130 135 140

Leu Pro Ala Asn Ile Thr Leu Lys Asp Val Pro Ser Leu Tyr Pro Ser 145 150 155 160

Phe His Ser Ala Ser Asp Ile Phe Asn Val Ala Lys Pro Lys Asn Pro 165 170 175

Ser Thr Asn Val Ser Val Val Val Phe Asp Ser Thr Lys Asp Val Glu 180 185 190

Asp Ala His Ser Gly Leu Leu Lys Gly Asn Ser Arg Gln Thr Val Trp 195 200 205

Arg Gly Tyr Leu Thr Thr Asp Lys Glu Val Pro Gly Leu Val Leu Met 210 215 220

Gln Asp Leu Ala Phe Leu Ser Gly Phe Pro Pro Thr Phe Lys Glu Thr 225 235 240

Asn Gln Leu Lys Thr Lys Leu Pro Glu Asn Leu Ser Ser Lys Val Lys 245 250 255

Leu Leu Gln Leu Tyr Ser Glu Ala Ser Val Ala Leu Leu Lys Leu Asn 260 265 270

Asn Pro Lys Asp Phe Gln Glu Leu Asn Lys Gln Thr Lys Lys Asn Met 275 280 285

Thr Ile Asp Gly Lys Glu Leu Thr Ile Ser Pro Ala Tyr Leu Leu Trp 290 295 300

Asp Leu Ser Ala Ile Ser Gln Ser Lys Gln Asp Glu Asp Ile Ser Ala 305 310 315 320

Ser Arg Phe Glu Asp Asn Glu Glu Leu Arg Tyr Ser Leu Arg Ser Ile 325 330 335

Glu Arg His Ala Pro Trp Val Arg Asn Ile Phe Ile Val Thr Asn Gly 340 345 350

Gln Ile Pro Ser Trp Leu Asn Leu Asp Asn Pro Arg Val Thr Ile Val

355 360 365

Thr His Gln Asp Val Phe Arg Asn Leu Ser His Leu Pro Thr Phe Ser 370 375 380

Ser Pro Ala Ile Glu Ser His Ile His Arg Ile Glu Gly Leu Ser Gln 385 390 395 400

Lys Phe Ile Tyr Leu Asn Asp Asp Val Met Phe Gly Lys Asp Val Trp 405 410 415

Pro Asp Asp Phe Tyr Ser His Ser Lys Gly Gln Lys Val Tyr Leu Thr 420 425 430

Trp Pro Val Pro Asn Cys Ala Glu Gly Cys Pro Gly Ser Trp Ile Lys
435 440 445

Asp Gly Tyr Cys Asp Lys Ala Cys Asn Asn Ser Ala Cys Asp Trp Asp 450 455 460

Gly Gly Asp Cys Ser Gly Asn Ser Gly Gly Ser Arg Tyr Ile Ala Gly 465 470 475 480

Gly Gly Gly Thr Gly Ser Ile Gly Val Gly His Pro Trp Gln Phe Gly 485 490 495

Gly Gly Ile Asn Ser Val Ser Tyr Cys Asn Gln Gly Cys Ala Asn Ser 500 505

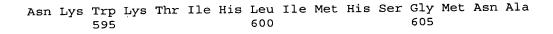
Trp Leu Ala Asp Lys Phe Cys Asp Gln Ala Cys Asn Val Leu Ser Cys 515 520 525

Gly Phe Asp Ala Gly Asp Cys Gly Gln Asp His Phe His Glu Leu Tyr 530 535 540

Lys Val Ile Leu Leu Pro Asn Gln Thr His Tyr Ile Ile Pro Lys Gly 545 550 555 560

Glu Cys Leu Pro Tyr Phe Ser Phe Ala Glu Val Ala Lys Arg Gly Val 565 570 575

Glu Gly Ala Tyr Ser Asp Asn Pro Ile Ile Arg His Ala Ser Ile Ala 580 585 590



Thr Thr Ile His Phe Asn Leu Thr Phe Gln Asn Thr Asn Asp Glu Glu 610 615 620

Phe Lys Met Gln Ile Thr Val Glu Val Asp Thr Arg Glu Gly Pro Lys 625 630 635

Leu Asn Ser Thr Ala Gln Lys Gly Tyr Glu Asn Leu Val Ser Pro Ile 645 650 655

Thr Leu Leu Pro Glu Ala Glu Ile Leu Phe Glu Asp Ile Pro Lys Glu 660 665 670

Lys Arg Phe Pro Lys Phe Lys Arg His Asp Val Asn Ser Thr Arg Arg 675 680 685

Ala Gln Glu Glu Val Lys Ile Pro Leu Val Asn Ile Ser Leu Leu Pro 690 695 700

Lys Asp Ala Gln Leu Ser Leu Asn Thr Leu Asp Leu Gln Leu Glu His 705 710 715 720

Gly Asp Ile Thr Leu Lys Gly Tyr Asn Leu Ser Lys Ser Ala Leu Leu 725 730 735

Arg Ser Phe Leu Met Asn Ser Gln His Ala Lys Ile Lys Asn Gln Ala 740 745 750

Ile Ile Thr Asp Glu Thr Asn Asp Ser Leu Val Ala Pro Gln Glu Lys 755 760 765

Gln Val His Lys Ser Ile Leu Pro Asn Ser Leu Gly Val Ser Glu Arg 770 775 780

Leu Gln Arg Leu Thr Phe Pro Ala Val Ser Val Lys Val Asn Gly His 785 790 795 800

Asp Gln Gly Gln Asn Pro Pro Leu Asp Leu Glu Thr Thr Ala Arg Phe 805 810 815 Arg Val Glu Thr His Thr Gln Lys Thr Ile Gly Gly Asn Val Thr Lys 820 825 830

Glu Lys Pro Pro Ser Leu Ile Val Pro Leu Glu Ser Gln Met Thr Lys 835 840 845

Glu Lys Lys Ile Thr Gly Lys Glu Lys Glu Asn Ser Arg Met Glu Glu 850 855 860

Asn Ala Glu Asn His Ile Gly Val Thr Glu Val Leu Leu Gly Arg Lys 865 870 875 880

Leu Gln His Tyr Thr Asp Ser Tyr Leu Gly Phe Leu Pro Trp Glu Lys 885 890 895

Lys Lys Tyr Phe Gln Asp Leu Leu Asp Glu Glu Glu Ser Leu Lys Thr 900 905 910

Gln Leu Ala Tyr Phe Thr Asp Ser Lys Asn Thr Gly Arg Gln Leu Lys 915 920 925

<210> 5

<211> 328

<212> PRT

<213> Homo sapiens

<400> 5

Asp Thr Phe Ala Asp Ser Leu Arg Tyr Val Asn Lys Ile Leu Asn Ser 1 5 10 15

Lys Phe Gly Phe Thr Ser Arg Lys Val Pro Ala His Met Pro His Met 20 25 30

Ile Asp Arg Ile Val Met Gl
n Glu Leu Gl
n Asp Met Phe Pro Glu Glu 35 40 45

Phe Asp Lys Thr Ser Phe His Lys Val Arg His Ser Glu Asp Met Gln 50 55 60

Phe Ala Phe Ser Tyr Phe Tyr Tyr Leu Met Ser Ala Val Gln Pro Leu 65 70 75 80

Asn Ile Ser Gln Val Phe Asp Glu Val Asp Thr Asp Gln Ser Gly Val

85 90 95

Leu Ser Asp Arg Glu Ile Arg Thr Leu Ala Thr Arg Ile His Glu Leu 100 105 110

Pro Leu Ser Leu Gln Asp Leu Thr Gly Leu Glu His Met Leu Ile Asn 115 120 125

Cys Ser Lys Met Leu Pro Ala Asp Ile Thr Gln Leu Asn Asn Ile Pro 130 135 140

Pro Thr Gln Glu Ser Tyr Tyr Asp Pro Asn Leu Pro Pro Val Thr Lys
145 150 155 160

Ser Leu Val Thr Asn Cys Lys Pro Val Thr Asp Lys Ile His Lys Ala 165 170 175

Tyr Lys Asp Lys Asn Lys Tyr Arg Phe Glu Ile Met Gly Glu Glu Glu 180 185 190

Ile Ala Phe Lys Met Ile Arg Thr Asn Val Ser His Val Val Gly Gln 195 200 205

Leu Asp Asp Ile Arg Lys Asn Pro Arg Lys Phe Val Cys Leu Asn Asp 210 215 220

Asn Ile Asp His Asn His Lys Asp Ala Gln Thr Val Lys Ala Val Leu 225 230 235 240

Arg Asp Phe Tyr Glu Ser Met Phe Pro Ile Pro Ser Gln Phe Glu Leu 245 250 255

Pro Arg Glu Tyr Arg Asn Arg Phe Leu His Met His Glu Leu Gln Glu 260 265 270

Trp Arg Ala Tyr Arg Asp Lys Leu Lys Phe Trp Thr His Cys Val Leu 275 280 285

Ala Thr Leu Ile Met Phe Thr Ile Phe Ser Phe Phe Ala Glu Gln Leu 290 295 300

Ile Ala Leu Lys Arg Lys Ile Phe Pro Arg Arg Arg Ile His Lys Glu 305 310 315 320

Ala Ser Pro Asn Arg Ile Arg Val 325

<210> 6 <211> 1219 <212> DNA <213> Homo sapiens <400> 6

60 gtagagegea ggtgegegge tegatggegg eggggetgge geggeteetg ttgeteeteg ggetetegge eggegggeee gegeeggeag gtgeagegaa gatgaaggtg gtggaggage 120 ccaacgcgtt tggggtgaac aacccgttct tgcctcaggc cagtcgcctc caggccaaga 180 gggatccttc accegtgtct ggaccegtgc atctcttccg actctcgggc aagtgcttca 240 gcctggtgga gtccacgtac aagtatgagt tctgcccgtt ccacaacgtg acccagcacg 300 360 agcagacett eegetggaae geetacagtg ggateetegg catetggeae gagtgggaga togocaacaa cacettoacg ggcatgtgga tgagggacgg tgacgcetge cgttcccgga 420 480 gccggcagag caaggtggag ctggcgtgtg gaaaaagcaa ccggctggcc catgtgtccg ageogageae etgegtetat gegetgaegt tegagaeece eetegtetge cacceccaeg 540 cettgetagt gtacceaace etgecagagg ceetgeageg geagtgggae caggtagage 600 660 aggacetgge egatgagetg ateaceecee agggeeatga gaagttgetg aggacaettt ttgaggatgc tggctactta aagaccccag aagaaaatga acccacccag ctggagggag 720 gtcctgacag cttggggttt gagaccctgg aaaactgcag gaaggctcat aaagaactct 780 840 caaaggagat caaaaggctg aaaggtttgc tcacccagca cggcatcccc tacacgaggc 900 ccacagaaac ttccaacttg gagcacttgg gccacgagac gcccagagcc aagtctccag agcagctgcg gggtgaccca ggactgcgtg ggagtttgtg accttgtggt gggagagcag 960 aggtggacgc ggccgagagc cctacagaga agctggctgg taggacccgc aggaccagct 1020 gaccaggett gtgeteagag aagcagacaa aacaaagatt caaggtttta attaatteee 1080 atactgataa aaataactcc atgaattctg taaaccattg cataaatgct atagtgtaaa 1140 1200 aaaatttaaa caagtgttaa ctttaaacag ttcgctacaa gtaaatgatt ataaatacta 1219 aaaaaaaaa aaaaaaaaa

<210> 7 <211> 305

<212> PRT

<213> Homo sapiens

<400> 7

Met Ala Ala Gly Leu Ala Arg Leu Leu Leu Leu Gly Leu Ser Ala 1 5 10 15

Gly Gly Pro Ala Pro Ala Gly Ala Ala Lys Met Lys Val Val Glu Glu 20 25 30

Pro Asn Ala Phe Gly Val Asn Asn Pro Phe Leu Pro Gln Ala Ser Arg 35 40 45

Leu Gln Ala Lys Arg Asp Pro Ser Pro Val Ser Gly Pro Val His Leu 50 55 60

Phe Arg Leu Ser Gly Lys Cys Phe Ser Leu Val Glu Ser Thr Tyr Lys 65 70 75 80

Tyr Glu Phe Cys Pro Phe His Asn Val Thr Gln His Glu Gln Thr Phe 85 90 95

Arg Trp Asn Ala Tyr Ser Gly Ile Leu Gly Ile Trp His Glu Trp Glu
100 105 110

Ile Ala Asn Asn Thr Phe Thr Gly Met Trp Met Arg Asp Gly Asp Ala 115 120 125

Cys Arg Ser Arg Ser Arg Gln. Ser Lys Val Glu Leu Ala Cys Gly Lys 130 135 140

Leu Thr Phe Glu Thr Pro Leu Val Cys His Pro His Ala Leu Leu Val 165 170 175

Tyr Pro Thr Leu Pro Glu Ala Leu Gln Arg Gln Trp Asp Gln Val Glu 180 185 190

Gln Asp Leu Ala Asp Glu Leu Ile Thr Pro Gln Gly His Glu Lys Leu 195 200 205

215 Asn Glu Pro Thr Gln Leu Glu Gly Gly Pro Asp Ser Leu Gly Phe Glu Thr Leu Glu Asn Cys Arg Lys Ala His Lys Glu Leu Ser Lys Glu Ile 250 Lys Arg Leu Lys Gly Leu Leu Thr Gln His Gly Ile Pro Tyr Thr Arg 260 265 Pro Thr Glu Thr Ser Asn Leu Glu His Leu Gly His Glu Thr Pro Arg 280 275 Ala Lys Ser Pro Glu Gln Leu Arg Gly Asp Pro Gly Leu Arg Gly Ser 300 Leu 305 <210> 8 <211> 5229 <212> DNA <213> Mus musculus <400> ggcggtgaag gggtgatgct gttcaagctc ctgcagagac agacctatac ctgcctatcc

Leu Arg Thr Leu Phe Glu Asp Ala Gly Tyr Leu Lys Thr Pro Glu Glu

60 cacaqqtatq qqctctacqt ctqcttcqtq qqcqtcqttq tcaccatcqt ctcqqctttc 120 cagttcggag aggtggttct ggaatggagc cgagatcagt accatgtttt gtttgattcc 180 tacagagaca acattgctgg gaaatccttt cagaatcggc tctgtctgcc catgccaatc 240 gacgtggttt acacctgggt gaatggcact gaccttgaac tgctaaagga gctacagcag 300 gtccgagagc acatggagga agagcagaga gccatgcggg aaaccctcgg gaagaacaca 360 accgaaccga caaagaagag tgagaagcag ctggaatgtc tgctgacgca ctgcattaag 420 gtgcccatgc ttgttctgga cccggccctg ccagccacca tcaccctgaa ggatctgcca 480 accetttace catettteca egegtecage gacatgttea atgttgegaa accaaaaaat 540 cegtetacaa atgteecegt tgtegttttt gacactacta aggatgttga agacgeecat 600 gctggaccgt ttaagggagg ccagcaaaca gatgtttgga gagcctactt gacaacagac 660 aaagacgccc ctggcttagt gctgatacaa ggcttggcgt tcctgagtgg attcccaccg 720 accttcaagg agacgagtca actgaagaca aagctgccaa gaaaagcttt ccctctaaaa 780 ataaagctgt tgcggctgta ctcggaggcc agtgtcgctc ttctgaaatt gaataatccc 840 aagggtttcc aagagctgaa caagcagacc aagaagaaca tgaccatcga tgggaaggaa 900 ctgaccatca gccctgcgta tctgctgtgg gacctgagtg ccatcagcca gtccaagcag 960 gatgaggacg cgtctgccag ccgctttgag gataatgaag agctgaggta ctcgctgcga 1020 1080 tctatcgaga gacacgcgcc atgggtacgg aatattttca ttgtcaccaa cgggcagatt 1140 ccatcctggc tgaaccttga caaccctcga gtgaccatag tgacccacca ggacattttc 1200 caaaatctga gccacttgcc tactttcagt tcccctgcta ttgaaagtca cattcaccgc 1260 atcgaagggc tgtcccagaa gtttatttat ctaaatgacg atgtcatgtt cggtaaggac gtctggccgg acgattttta cagccactcc aaaggtcaaa aggtttattt gacatggcct 1320 1380 gtgccaaact gtgcagaggg ctgcccgggc tcctggataa aggacggcta ttgtgataag gcctgtaata cctcaccctg tgactgggat ggcggaaact gctctggtaa tactgcaggg 1440 1500 aaccggtttg ttgcaagagg tgggggtacc gggaatattg gagctggaca gcactggcag 1560 tttggtggag gaataaacac catctcttac tgtaaccaag gatgtgcaaa ctcctggctg 1620 gctgacaagt tctgtgacca agcctgtaac gtcttatcct gcgggtttga tgctggtgac tgtggacaag atcattttca tgaattgtat aaagtaacac ttctcccaaa ccagactcac 1680 tatgttgtcc ccaaaggtga atacctgtct tatttcagct ttgcaaacat agccagaaaa 1740 1800 agaattgaag ggacctacag cgacaacccc atcatccgcc acgcgtccat tgcaaacaag 1860 tggaaaaccc tacacctgat aatgcccggg gggatgaacg ccaccacgat ctattttaac 1920 ctcactcttc aaaacgccaa cgacgaagag ttcaagatcc agatagcagt agaggtggac 1980 acgagggagg cgcccaaact gaattctaca acccagaagg cctatgaaag tttggttagc 2040 ccagtgacac ctcttcctca ggctgacgtc ccttttgaag atgtccccaa agagaaacgc 2100 ttccccaaga tcaggagaca tgatgtaaat gcaacaggga gattccaaga ggaggtgaaa atccccggg taaatatttc actccttccc aaagaggccc aggtgaggct gagcaacttg 2160 gatttgcaac tagaacgtgg agacatcact ctgaaaggat ataacttgtc caagtcagcc 2220 ctgctaaggt ctttcctggg gaattcacta gatactaaaa taaaacctca agctaggacc 2280 gatgaaacaa aaggcaacct ggaggtccca caggaaaacc cttctcacag acgtccacat 2340 2400 ggctttgctg qtgaacacag atcagagaga tggactgccc cagcagagac agtgaccgtg

aaaggeegtg accaegettt gaateeacee eeggtgttgg agaeeaatge aagattggee 2460 cagcctacac taggcgtgac tgtgtccaaa gagaaccttt caccgctgat cgttccccca 2520 gaaagccact tgccaaaaga agaggagagt gacagggcag aaggcaatgc tgtacctgta 2580 aaggagttag tgcctggcag acggttgcag cagaattatc caggcttttt gccctgggag 2640 aaaaaaagt atttccaaga ccttcttgat gaggaagagt cattgaagac ccagttggcg 2700 tactttacag accgcaaaca taccgggagg caactaaaag atacatttgc agactccctc 2760 cgatacgtca ataaaattct caacagcaag tttggattca catccaggaa agtccctgca 2820 cacatgccgc acatgattga caggatcgtt atgcaagaac tccaagatat gttccctgaa 2880 gaatttgaca agacttcatt tcacaaggtg cgtcactctg aggacatgca gtttgccttc 2940 tectaetttt attaeeteat gagtgeagtt eageceetea atattteeea agtettteat 3000 gaagtagaca cagaccaatc tggtgtcttg tctgataggg aaatccgaac wctggccacg 3060 agaattcacg acctaccttt aagcttgcag gatttgacag gtttggaaca catgttaata 3120 3180 gaagcatact acgaccccaa cctgcctccg gtcactaaga gtcttgtcac caactgtaag 3240 ccagtaactg acaagatcca caaagcctat aaagacaaga acaaatacag gtttgaaatc 3300 atgggagagg aagaaatcgc tttcaagatg atacgaacca atgtttctca tgtggttggt 3360 cagttggatg acatcagaaa aaaccccagg aagttcgttt gtctgaatga caacattgac 3420 cacaaccata aagatgcccg gacagtgaag gctgtcctca gggacttcta tgagtccatg 3480 tttcccatac cttcccagtt tgagctgcca agagagtatc ggaaccgctt tctgcacatg 3540 catgagetee aagaatggeg ggeatatega gacaagetga agttttggae eeactgegta 3600 ctagcaacgt tgattatatt tactatattc tcattttttg ctgaacagat aattgctctg 3660 aagcgaaaga tatttcccag gaggaggata cacaaagaag ctagtccaga ccgaatcagg 3720 gtgtagaaga tottcatttg aaagtcacct accttagcat ctgtgaacat ctccctcctc 3780 gacaccacag eggagteect gtgatgtgge acagaggeag cetegtgggg agaagggaca 3840 tegtgeagae egggttette tgeaatggga agagageeea etgaeetgga attatteage 3900 acactaagaa cctgtgtcaa tagcttgtac agcttgtact tttaaaggat ttgccgaagg 3960 acctgtcggc ttgttgacaa accctccctg acaagctgct ggtttcttcc cccagttact 4020 gcagactgag aaaccagtcc atcttgaaag caagtgcgga ggggccccag tctttgcatt 4080 ccaaagettt ccagcataat ttctggettg teteeteett tgateeattt cccatttttt 4140

| tttaaaaaac | aataagtggc | tactaagtta | gtcattctca | cttctcaaaa | taacaaatca | 4200 |
|------------|------------|------------|------------|------------|------------|------|
| ggatgtcaaa | acatttgtat | agatcttatt | taaataatat | agaacgatta | cttctttagc | 4260 |
| ctatctaaat | tattgatttt | tattaacagt | caagtggtct | tgaaccgcta | acaactactg | 4320 |
| aagagctcga | gattgacgtt | gaaagtgctt | tgagcttgtt | taactcattc | cccaagaata | 4380 |
| ctgtgacctc | gtgtgcgggc | ctgattgcga | agggctagtg | tcacgtagca | gtgctgctca | 4440 |
| ccggatgtaa | ttatgtcgtg | gaaatgtaca | tacagacaaa | agtgcctcac | ttcagaaatg | 4500 |
| agtagtgctg | atggcaccag | cgagtgatgg | tgtccatttg | gaaacccatg | ataccttcca | 4560 |
| atgcccaccc | tgcttacttt | atacagagca | ggggttaacc | aacttctgtc | aaagaacagt | 4620 |
| aaagaacttg | agatacatcc | atctttgtca | aatagttttc | cttgctaaca | tttattattg | 4680 |
| ttggtgtttt | gggaggttta | ttttatttta | ttgctttgtt | atttttcaag | acggggattc | 4740 |
| tctgtgtagc | tctggctgtt | tggtaattca | ctctaaagac | caggctggcc | ttgaacttag | 4800 |
| agattcacct | gcttctgctt | cctgaatggt | aggacatgtg | cccacattgc | ctacccaccc | 4860 |
| cccttttggg | gggggtgagc | aactcaataa | aaagatgaaa | acctgcttta | gtttgcagct | 4920 |
| atacaaaagc | agcaggcctc | agccagactt | gacccccggg | gccattgttg | gcccacggga | 4980 |
| gaatcatttt | tgacgtgggt | aagcaaaccc | tgatattggt | catgctgtgt | tatgtcatta | 5040 |
| tgtggtggtt | ttgaattttg | gaagatattt | tcagtcatga | tttcagtagt | attcctccaa | 5100 |
| aatggcacac | atttttgtaa | taagaacttg | aaatgtaaat | attgtgtttg | tgctgtaaat | 5160 |
| tttgtgtatt | tcaaaaactg | aagtttcata | aaaaaacaca | cttattggaa | aaaaaaaaa | 5220 |
| aaaaaaaaa | | | | | | 5229 |

<210> 9

Met Leu Phe Lys Leu Leu Gln Arg Gln Thr Tyr Thr Cys Leu Ser His 10

Arg Tyr Gly Leu Tyr Val Cys Phe Val Gly Val Val Val Thr Ile Val

Ser Ala Phe Gln Phe Gly Glu Val Val Leu Glu Trp Ser Arg Asp Gln . 40

<211> 908 <212> PRT <213> Mus musculus

<400> 9

Tyr His Val Leu Phe Asp Ser Tyr Arg Asp Asn Ile Ala Gly Lys Ser 50 55 60

Phe Gln Asn Arg Leu Cys Leu Pro Met Pro Ile Asp Val Val Tyr Thr 65 70 75 80

Trp Val Asn Gly Thr Asp Leu Glu Leu Leu Lys Glu Leu Gln Gln Val 85 90 95

Arg Glu His Met Glu Glu Glu Gln Arg Ala Met Arg Glu Thr Leu Gly 100 105 110

Lys Asn Thr Thr Glu Pro Thr Lys Lys Ser Glu Lys Gln Leu Glu Cys 115 120 125

Leu Leu Thr His Cys Ile Lys Val Pro Met Leu Val Leu Asp Pro Ala 130 135 140

Leu Pro Ala Thr Ile Thr Leu Lys Asp Leu Pro Thr Leu Tyr Pro Ser 145 150 155 160

Phe His Ala Ser Ser Asp Met Phe Asn Val Ala Lys Pro Lys Asn Pro 165 170 175

Ser Thr Asn Val Pro Val Val Val Phe Asp Thr Thr Lys Asp Val Glu 180 185 190

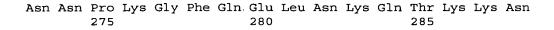
Asp Ala His Ala Gly Pro Phe Lys Gly Gly Gln Gln Thr Asp Val Trp 195 200 205

Arg Ala Tyr Leu Thr Thr Asp Lys Asp Ala Pro Gly Leu Val Leu Ile 210 215 220

Gln Gly Leu Ala Phe Leu Ser Gly Phe Pro Pro Thr Phe Lys Glu Thr 225 230 235 240

Ser Gln Leu Lys Thr Lys Leu Pro Arg Lys Ala Phe Pro Leu Lys Ile 245 250 255

Lys Leu Leu Arg Leu Tyr Ser Glu Ala Ser Val Ala Leu Leu Lys Leu 260 265 270



Met Thr Ile Asp Gly Lys Glu Leu Thr Ile Ser Pro Ala Tyr Leu Leu 290 295 300

Trp Asp Leu Ser Ala Ile Ser Gln Ser Lys Gln Asp Glu Asp Ala Ser 305 310 315 320

Ala Ser Arg Phe Glu Asp Asn Glu Glu Leu Arg Tyr Ser Leu Arg Ser 325 330 335

Ile Glu Arg His Ala Pro Trp Val Arg Asn Ile Phe Ile Val Thr Asn 340 345 350

Gly Gln Ile Pro Ser Trp Leu Asn Leu Asp Asn Pro Arg Val Thr Ile 355 360 365

Val Thr His Gln Asp Ile Phe Gln Asn Leu Ser His Leu Pro Thr Phe 370 375 380

Ser Ser Pro Ala Ile Glu Ser His Ile His Arg Ile Glu Gly Leu Ser 385 390 395 400

Gln Lys Phe Ile Tyr Leu Asn Asp Asp Val Met Phe Gly Lys Asp Val
405 410 415

Trp Pro Asp Asp Phe Tyr Ser His Ser Lys Gly Gln Lys Val Tyr Leu 420 425 430

Thr Trp Pro Val Pro Asn Cys Ala Glu Gly Cys Pro Gly Ser Trp Ile 435 440 445

Lys Asp Gly Tyr Cys Asp Lys Ala Cys Asn Thr Ser Pro Cys Asp Trp 450 455 460

Asp Gly Gly Asn Cys Ser Gly Asn Thr Ala Gly Asn Arg Phe Val Ala 465 470 475 480

Arg Gly Gly Gly Thr Gly Asn Ile Gly Ala Gly Gln His Trp Gln Phe 485 490 495 Gly Gly Gly Ile Asn Thr Ile Ser Tyr Cys Asn Gln Gly Cys Ala Asn 500 505

Ser Trp Leu Ala Asp Lys Phe Cys Asp Gln Ala Cys Asn Val Leu Ser 515 520 525

Cys Gly Phe Asp Ala Gly Asp Cys Gly Gln Asp His Phe His Glu Leu 530 535 540

Tyr Lys Val Thr Leu Leu Pro Asn Gln Thr His Tyr Val Val Pro Lys 545 550 560

Gly Glu Tyr Leu Ser Tyr Phe Ser Phe Ala Asn Ile Ala Arg Lys Arg 565 570 575

Ile Glu Gly Thr Tyr Ser Asp Asn Pro Ile Ile Arg His Ala Ser Ile 580 585

Ala Asn Lys Trp Lys Thr Leu His Leu Ile Met Pro Gly Gly Met Asn 595 600 605

Ala Thr Thr Ile Tyr Phe Asn Leu Thr Leu Gln Asn Ala Asn Asp Glu 610 620

Glu Phe Lys Ile Gln Ile Ala Val Glu Val Asp Thr Arg Glu Ala Pro 625 630 635 640

Lys Leu Asn Ser Thr Thr Gln Lys Ala Tyr Glu Ser Leu Val Ser Pro 645 650 655

Val Thr Pro Leu Pro Gln Ala Asp Val Pro Phe Glu Asp Val Pro Lys 660 665 670

Glu Lys Arg Phe Pro Lys Ile Arg Arg His Asp Val Asn Ala Thr Gly 675 680 685

Arg Phe Gln Glu Glu Val Lys Ile Pro Arg Val Asn Ile Ser Leu Leu 690 695 700

Pro Lys Glu Ala Gln Val Arg Leu Ser Asn Leu Asp Leu Gln Leu Glu 705 710 715 720

Arg Gly Asp Ile Thr Leu Lys Gly Tyr Asn Leu Ser Lys Ser Ala Leu

725 730 735

Leu Arg Ser Phe Leu Gly Asn Ser Leu Asp Thr Lys Ile Lys Pro Gln
740 745 750

Ala Arg Thr Asp Glu Thr Lys Gly Asn Leu Glu Val Pro Gln Glu Asn 755 760 765

Pro Ser His Arg Arg Pro His Gly Phe Ala Gly Glu His Arg Ser Glu 770 780

Arg Trp Thr Ala Pro Ala Glu Thr Val Thr Val Lys Gly Arg Asp His 785 790 795 800

Ala Leu Asn Pro Pro Pro Val Leu Glu Thr Asn Ala Arg Leu Ala Gln 805 810 815

Pro Thr Leu Gly Val Thr Val Ser Lys Glu Asn Leu Ser Pro Leu Ile 820 825 830

Val Pro Pro Glu Ser His Leu Pro Lys Glu Glu Glu Ser Asp Arg Ala 835 840 845

Glu Gly Asn Ala Val Pro Val Lys Glu Leu Val Pro Gly Arg Arg Leu 850 855 860

Gln Gln Asn Tyr Pro Gly Phe Leu Pro Trp Glu Lys Lys Lys Tyr Phe 865 870 875

Gln Asp Leu Leu Asp Glu Glu Glu Ser Leu Lys Thr Gln Leu Ala Tyr 885 890 895

Phe Thr Asp Arg Lys His Thr Gly Arg Gln Leu Lys 900 905

<210> 10

<211> 328

<211> 328 <212> PRT

<213> Mus musculus

<400> 10

Asp Thr Phe Ala Asp Ser Leu Arg Tyr Val Asn Lys Ile Leu Asn Ser 1 5 10 15

Lys Phe Gly Phe Thr Ser Arg Lys Val Pro Ala His Met Pro His Met 20 25 30

Ile Asp Arg Ile Val Met Gln Glu Leu Gln Asp Met Phe Pro Glu Glu 35 40 45

Phe Asp Lys Thr Ser Phe His Lys Val Arg His Ser Glu Asp Met Gln 50 55 60

Phe Ala Phe Ser Tyr Phe Tyr Tyr Leu Met Ser Ala Val Gln Pro Leu 65 70 75 80

Asn Ile Ser Gln Val Phe His Glu Val Asp Thr Asp Gln Ser Gly Val 85 90 95

Leu Ser Asp Arg Glu Ile Arg Thr Leu Ala Thr Arg Ile His Asp Leu 100 105 110

Pro Leu Ser Leu Gln Asp Leu Thr Gly Leu Glu His Met Leu Ile Asn 115 120 125

Cys Ser Lys Met Leu Pro Ala Asn Ile Thr Gln Leu Asn Asn Ile Pro 130 135 140

Pro Thr Gln Glu Ala Tyr Tyr Asp Pro Asn Leu Pro Pro Val Thr Lys 145 150 155 160

Ser Leu Val Thr Asn Cys Lys Pro Val Thr Asp Lys Ile His Lys Ala 165 170 175

Tyr Lys Asp Lys Asn Lys Tyr Arg Phe Glu Ile Met Gly Glu Glu Glu 180 185 190

Ile Ala Phe Lys Met Ile Arg Thr Asn Val Ser His Val Val Gly Gln
195 200 205

Leu Asp Asp Ile Arg Lys Asn Pro Arg Lys Phe Val Cys Leu Asn Asp 210 215 220

Asn Ile Asp His Asn His Lys Asp Ala Arg Thr Val Lys Ala Val Leu 225 230 235 240

| Arg | Asp | Phe | Tyr | Glu | Ser | Met | Phe | Pro | Ile | Pro | Ser | Gln | Phe | Glu | Leu |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| | | | | 245 | | • | | | 250 | | | | | 255 | |

Pro Arg Glu Tyr Arg Asn Arg Phe Leu His Met His Glu Leu Gln Glu

Trp Arg Ala Tyr Arg Asp Lys Leu Lys Phe Trp Thr His Cys Val Leu 280

Ala Thr Leu Ile Ile Phe Thr Ile Phe Ser Phe Phe Ala Glu Gln Ile

Ile Ala Leu Lys Arg Lys Ile Phe Pro Arg Arg Ile His Lys Glu 310 315

Ala Ser Pro Asp Arg Ile Arg Val 325

<210> 11 <211> 2070 <212> 'DNA <213> Mus musculus <220> <221> misc_feature

(186)..(186) <223> n is a, t, c, or g

<400> 11

<222>

gtgagaccet aggageaatg geegggegge tggetggett eetgatgttg etggggeteg cgtcgcaggg gcccgcgccg gcatgtgccg ggaagatgaa ggtggtggag gagcctaaca cattegggtg ageggateae ggteetgegg ettggggace gageetgget ggttettetg accttntcaa ttccataggc tgaataaccc gttcttgccc caggcaagcc gccttcagcc caagagagag ccttcagctg tatcccgcaa attaagagaa attaatttca aacgatttag aaagtattot agocaggoga tgatggogoa ogootttaat oocagcactt gggaggoaga ggcaggcaga tttccgagtt caaggccatc agaactgact gtacatctta gtacagttta gcatgtgatc agagatctga atcacaaagc tgggcctgcg tggtaaagca ggtcctttct aataaggttg cagtttagat tttctttctt aactctttta ttctttgaga cagggtttct caacagtggg tgtcctggaa ctcacttttg taaaccaggc tgcccttaaa ctcacaaagc tetgteagee tetgeeteet gagtgetggg attaaaggte caeaccetgt teatteattt 60

120

180

240

300

360

420

480

540

600

660

| ttaatttttg | agactgggtc | tcattatgtg | gccctagaca | gatactgaga | gcctcctcca | 720 |
|------------|------------|------------|------------|------------|------------|------|
| caggaacaag | catgggaatc | ctgccacaga | caaccagttc | tgtggtctgg | agatgagttt | 780 |
| gtcagtccct | aggagttagg | tcagcctgcc | tctgcattcc | caataattta | ggaaaggagc | 840 |
| ttggggcgtt | ctggccttga | tggttagtgc | cctcctgcca | accttagctt | ccagctttag | 900 |
| gggtagcaga | gtttataccg | atgctaaact | gctgttgtgt | tcttccccag | ggcccctgca | 960 |
| tctcttcaga | cttgctggca | agtgctttag | cctagtggag | tccacgtgag | tgccaggctg | 1020 |
| gtgggtggag | tgggcggagt | ctgcagagct | cctgatgtgc | ctgtgtttcc | caggtacaag | 1080 |
| tatgaattct | gccctttcca | caacgtcacc | cagcacgagc | agaccttccg | ctggaatgcc | 1140 |
| tacagcggga | tccttggcat | ctggcatgag | tgggaaatca | tcaacaatac | cttcaagggc | 1200 |
| atgtggatga | ctgatgggga | ctcctgccac | teceggagee | ggcagagcaa | ggtggagctc | 1260 |
| acctgtggaa | agatcaaccg | actggcccac | gtgtctgagc | caagcacctg | tgtctatgca | 1320 |
| ttgacattcg | agacccctct | tgtttgccat | cccactctt | tgttagtgta | tccaactctg | 1380 |
| tcagaagccc | tgcagcagcc | cttggaccag | gtggaacagg | acctggcaga | tgaactgatc | 1440 |
| acaccacagg | gctatgagaa | gttgctaagg | gtactttttg | aggatgctgg | ctacttaaag | 1500 |
| gtcccaggag | aaacccatcc | cacccagctg | gcaggaggtt | ccaagggcct | ggggcttgag | 1560 |
| actctggaca | actgtagaaa | ggcacatgca | gagctgtcac | aggaggtaca | aagactgacg | 1620 |
| agtctgctgc | aacagcatgg | aatcccccac | actcagccca | caggtcagtc | tgcctgccct | 1680 |
| ggtcagctgc | cagccactcc | ggggcctgca | gcactggggc | agatctttat | tgctacccat | 1740 |
| tctggcagaa | accactcact | ctcagcacct | gggtcagcag | ctccccatag | gtgcaatcgc | 1800 |
| agcagagcat | ctgcggagtg | acccaggact | acgtgggaac | atcctgtgag | caaggtggcc | 1860 |
| acgaagaata | gaaatatcct | gagctttgag | tgtcctttca | cagagtgaac | aaaactggtg | 1920 |
| tggtgtagac | acggcttctt | ttggcatatt | ctagatcaga | cagtgtcact | gacaaacaag | 1980 |
| agggacctgc | tggccagcct | ttgttgtgcc | caaagatcca | gacaaaataa | agattcaaag | 2040 |
| ttttaattaa | aaaaaaaaa | aaaggaatto | ! | | | 2070 |

<210> 12 <211> 307

<212> PRT <213> Mus musculus

<400> 12

Met Ala Gly Arg Leu Ala Gly Phe Leu Met Leu Leu Gly Leu Ala Ser 1 5 10 15

Gln Gly Pro Ala Pro Ala Cys Ala Gly Lys Met Lys Val Val Glu Glu 20 25 30

Pro Asn Thr Phe Gly Leu Asn Asn Pro Phe Leu Pro Gln Ala Ser Arg 35 40 45

Leu Gln Pro Lys Arg Glu Pro Ser Ala Val Ser Gly Pro Leu His Leu 50 55 60

Phe Arg Leu Ala Gly Lys Cys Phe Ser Leu Val Glu Ser Thr Tyr Lys 65 70 75 80

Tyr Glu Phe Cys Pro Phe His Asn Val Thr Gln His Glu Gln Thr Phe 85 90 95

Arg Trp Asn Ala Tyr Ser Gly Ile Leu Gly Ile Trp His Glu Trp Glu
100 105 110

Ile Ile Asn Asn Thr Phe Lys Gly Met Trp Met Thr Asp Gly Asp Ser 115 120 125

Cys His Ser Arg Ser Arg Gln Ser Lys Val Glu Leu Thr Cys Gly Lys 130 135 140

Leu Thr Phe Glu Thr Pro Leu Val Cys His Pro His Ser Leu Leu Val 165 170 175

Tyr Pro Thr Leu Ser Glu Ala Leu Gln Gln Arg Leu Asp Gln Val Glu 180 . 185 190

Gln Asp Leu Ala Asp Glu Leu Ile Thr Pro Gln Gly Tyr Glu Lys Leu 195 200 205

Leu Arg Val Leu Phe Glu Asp Ala Gly Tyr Leu Lys Val Pro Gly Glu 210 215 220

Thr His Pro Thr Gln Leu Ala Gly Gly Ser Lys Gly Leu Gly Leu Glu

| 1111111 | _ | c |
|---------|-------|----------|
| Ę | = | 3 |
| į | = | ¥ |
| F | Ti. | Į. |
| Ę | į | ļ |
| į | ĭ | in the |
| | Ĭ | |
| Ī | Į | Ţ |
| E | | |
| E_ | = | į. |
| F | ī, | i i |
| 2 | and a | three. |
| Ļ | = | Ŀ |
| Ē | = | ş |
| _ | | <u>.</u> |

<400> 14

| Thr I | Leu | Asp | Asn | Cys 245 | Arg | Lys | Ala | His | Ala 250 | Glu | Leu | Ser | Gln | Glu 255 | Val | |
|----------------------------------|------------|--------------------------|------------|------------|-------|------------------|------------|------------|------------|------|------------|------------|------------|------------|-------|-------|
| Gln A | Arg | Leu | Thr 260 | Ser | Leu | Leu _. | Gln | Gln 265 | His | Gly | Ile | Pro | His 270 | Thr | Gln | |
| Pro T | Thr | Glu 275 | Thr | Thr | His | Ser | Gln 280 | His | Leu | Gly | Gln | Gln 285 | Leu | Pro | Ile | |
| Gly A | Ala 290 | Ile | Ala | Ala | Glu | His 295 | Leu | Arg | Ser | Asp | Pro 300 | Gly | Leu | Arg | Gly | |
| Asn 3 | Ile | Leu | | | | | | | | | | | | | | |
| <210: <211: <212: <213: | > 4 > 1 | 13 460 DNA Ratt | us r | attu | s | | | | | | | | | | | |
| <400 | | 13 cca | acat | tcaa | gg a | gacg | agtc | a gc | tgaa | gaca | aaa | ctgc | cag | aaaa | tcttt | c 60 |
| | | | | | | | | | | | | | | | attga | |
| | | | | | | | | | | | | | | | cagtg | |
| gaag | gaa | ctg | gcca | tcag | cc c | tgcc | tato | t go | tgtg | ggac | ctg | agcg | cca | tcag | ccagt | c 240 |
| caag | ıcag | gat | gaag | atgt | gt c | tgcc | agco | g ct | tcga | ggat | aac | gaag | agc | tgag | gtact | c 300 |
| actg | jaga | tct | atcg | agag | ac a | tgat | tcca | it ga | gtco | ttta | . tga | atto | tgg | ccat | atctt | c 360 |
| aatc | atg | atc | tcag | tagt | at t | cctc | tgaa | a te | gcac | acat | ttt | tcta | atg | agaa | cttga | a 420 |
| atgt | aaa | tat | tgtg | ıtttg | ıtg c | tgta | aatt | t te | gtgta | tttc | : | | | | | 460 |
| <210 <211 <212 <213 | L> 2> | 14 113 PRT Ratt | us 1 | ratti | ıs | | | | | | | | | | | |

Phe Pro Pro Thr Phe Lys Glu Thr Ser Gln Leu Lys Thr Lys Leu Pro

Glu Asn Leu Ser Ser Lys Ile Lys Leu Leu Gln Leu Tyr Ser Glu Ala

Ser Val Ala Leu Leu Lys Leu Asn Asn Pro Lys Gly Phe Pro Glu Leu 40

Asn Lys Gln Thr Lys Lys Asn Met Ser Ile Ser Gly Lys Glu Leu Ala

Ile Ser Pro Ala Tyr Leu Leu Trp Asp Leu Ser Ala Ile Ser Gln Ser 70

Lys Gln Asp Glu Asp Val Ser Ala Ser Arg Phe Glu Asp Asn Glu Glu

Leu Arg Tyr Ser Leu Arg Ser Ile Glu Arg His Asp Ser Met Ser Pro 105

Leu

<210> 15

<211> 1105

<212> DNA

<213> Drosophila melanogaster

<220>

<221> misc_feature

(903)..(903) <222>

<223> n is a, g, t, or c

<220>

<221> misc feature

<222> (935)..(935)

<223> n is a, g, t, or c

<220>

<221> misc_feature

<222> (1023)..(1023) <223> n is a, g, t, or c

<220>

<221> misc_feature

<222> (1035)..(1035)

<223> n is a, g, t, or c

```
<220>
      misc feature
<221>
      (1071)..(1071)
<222>
<223> n is a, g, t, or c
<220>
      misc feature
<221>
       (1100)..(1100)
<222>
      n is a, g, t, or c
<400> 15
ctgcaggaat tcggcacgag gcggttcgat gacaagaatg agctgcggta ctctctgagg
                                                                       60
tccctggaaa aacacgccgc atggatcagg catgtgtaca tagtaaccaa tggccagatt
                                                                      120
ccaagttggc tggatctcag ctacgaaagg gtcacggtgg tgccccacga agtcctggct
                                                                      180
cccgatcccg accagctgcc caccttctcc agetcggcca tcgagacatt tctgcaccgc
                                                                      240
ataccaaagc tgtccaagag gttcctctac ctcaacgacg acatattcct gggagctccg
                                                                      300
ctgtatccgg aggacttgta cactgaagcg gagggagttc gcgtgtacca ggcatggatg
                                                                       360
gtgcccggct gcgccttgga ttgcccctgg acgtacatag gtgatggagc ttgcgatcgg
                                                                       420
cactgcaaca ttgatgcgtg ccaatttgat ggaggcgact gcagtgaaac tgggccagcg
                                                                       480
agegatgeec aegteattee accaageaaa gaagtgeteg aggtgeagee tgeegetgtt
                                                                       540
ccacaatcaa gagtccaccg atttcctcag atgggtctcc aaaagctgtt caggcgcagc
                                                                       600
 tctgccaatt ttaaggatgt tatgcggcac cgcaatgtgt ccacactcaa ggaactacgt
                                                                       660
 cgcattgtgg agcgttttaa caaggccaaa ctcatgtcgc tgaaccccga actggagacc
                                                                       720
                                                                       780
 tccagctccg agccacagac aactcagcgc cacgggctgc gcaaggagga ttttaagtct
```

tecacegata tttaetetea etegetgatt gecaceaata tgttgetgaa tagageetat

ggctttaagg cacgccatgt cctggcgcac gtgggcttcc taattgacaa ggatattgtg

gangccatgc aacgacgttt taccagcgaa ttctngacac tggccattaa cgctttccga

gccccaacag atttgcagta cgcattcgct tactacttct ttctaatgag cgaaatccaa

gtnatgagtg tagangaaat cttcgatgaa gtcgacaccg gacggtttgg ncacctggtc

ggatccagaa gtgcgaaccn tttta

840

900

960

1020

1080

1105

<210> 16 <211> 502 <212> PRT <213> Drosophila melanogaster

<400> 16

Gly Thr Arg Arg Phe Asp Asp Lys Asn Glu Leu Arg Tyr Ser Leu Arg 1 5 10 15

Ser Leu Glu Lys His Ala Ala Trp Ile Arg His Val Tyr Ile Val Thr 20 25 30

Asn Gly Gln Ile Pro Ser Trp Leu Asp Leu Ser Tyr Glu Arg Val Thr 35 40 45

Val Val Pro His Glu Val Leu Ala Pro Asp Pro Asp Gln Leu Pro Thr 50 60

Phe Ser Ser Ser Ala Ile Glu Thr Phe Leu His Arg Ile Pro Lys Leu 65 70 75 80

Ser Lys Arg Phe Leu Tyr Leu Asn Asp Asp Ile Phe Leu Gly Ala Pro 85 90 95

Leu Tyr Pro Glu Asp Leu Tyr Thr Glu Ala Glu Gly Val Arg Val Tyr 100 105 110

Gln Ala Trp Met Val Pro Gly Cys Ala Leu Asp Cys Pro Trp Thr Tyr 115 120 125

Ile Gly Asp Gly Ala Cys Asp Arg His Cys Asn Ile Asp Ala Cys Gln 130 135 140

Phe Asp Gly Gly Asp Cys Ser Glu Thr Gly Pro Ala Ser Asp Ala His 145 150 155 160

Val Ile Pro Pro Ser Lys Glu Val Leu Glu Val Gln Pro Ala Ala Val 165 170 175

Pro Gln Ser Arg Val His Arg Phe Pro Gln Met Gly Leu Gln Lys Leu 180 185 190

Phe Arg Arg Ser Ser Ala Asn Phe Lys Asp Val Met Arg His Arg Asn 195 200 205

Val Ser Thr Leu Lys Glu Leu Arg Arg Ile Val Glu Arg Phe Asn Lys 210 215 220

Ala Lys Leu Met Ser Leu Asn Pro Glu Leu Glu Thr Ser Ser Ser Glu 240

Pro Gln Thr Thr Gln Arg His Gly Leu Arg Lys Glu Asp Phe Lys Ser 255

Ser Thr Asp Ile Tyr Ser His Ser Leu Ile Ala Thr Asn Met Leu Leu 260 265 270

Asn Arg Ala Tyr Gly Phe Lys Ala Arg His Val Leu Ala His Val Gly 275 280 285

Phe Leu Ile Asp Lys Asp Ile Val Glu Ala Met Gln Arg Arg Phe His 290 295 300

Gln Gln Ile Leu Asp Thr Ala His Gln Arg Phe Arg Ala Pro Thr Asp 305 310 315 320

Leu Gln Tyr Ala Phe Ala Tyr Tyr Ser Phe Leu Met Ser Glu Thr Lys 325 330 335

Val Met Ser Val Glu Glu Ile Phe Asp Glu Phe Asp Thr Asp Gly Ser 340 345 350

Ala Thr Trp Ser Asp Arg Glu Val Arg Thr Phe Leu Thr Arg Ile Tyr 355 360 365

Gln Pro Pro Leu Asp Trp Ser Ala Met Arg Tyr Phe Glu Glu Val Val 370 375 380

Gln Asn Cys Thr Arg Asn Leu Gly Met His Leu Lys Val Asp Thr Val 385 390 395 400

Glu His Ser Thr Leu Val Tyr Glu Arg Tyr Glu Asp Ser Asn Leu Pro 405 410 415

Thr Ile Thr Arg Asp Leu Val Val Arg Cys Pro Leu Leu Ala Glu Ala 420 425 430

Leu Ala Ala Asn Phe Ala Val Arg Pro Lys Tyr Asn Phe His Val Ser 435 .440 .445 Pro Lys Arg Thr Ser His Ser Asn Phe Met Met Leu Thr Ser Asn Leu 450 455 460

Thr Glu Val Val Glu Ser Leu Asp Arg Leu Arg Arg Asn Pro Arg Lys
465 470 475 480

Phe Asn Cys Ile Asn Asp Asn Leu Asp Ala Asn Arg Gly Glu Asp Asn 490 495

Glu Asp Gly Ala Pro Ser 500

<210> 17 <211> 2183 <212> DNA

<213> Homo sapiens

<400> 17 60 atggcgacct ccacgggtcg ctggcttctc ctccggcttg cactattcgg cttcctctgg gaagegteeg geggeetega etegggggee teeegegaeg aegaettget aetgeeetat 120 ccacgcgcgc gegcgcct cccccgggac tgcacacggg tgcgcccgg caaccgcgag 180 cacgagagtt ggcctccgcc tcccgcgact cccggcgccg gcggtctggc cgtgcgcacc 240 300 ttoqtqtoqo acttoaggga cogogoggtg googgocaco tgaogogggo ogttgagoco ctgcgcacct tctcggtgct ggagcccggt ggacccggcg gctgcgcggc gagacgacgc 360 420 gccaccgtgg aggagacggc gcgggcggcc gactgccgtg tcgcccagaa cggcggcttc ttccqcatga actcgggcga gtgcctgggg aacgtggtga gcgacgagcg gcgggtgagc 480 ageteegggg ggetgeagaa egegeagtte gggateegee gegaegggae eetggteace 540 gggtacctgt ctgaggagga ggtgctggac actgagaacc catttgtgca gctgctgagt 600 660 ggggtcgtgt ggctgattcg taatggaagc atctacatca acgagagcca agccacagag tgtgacgaga cacaggagac aggttccttt agcaaatttg tgaatgtgat atcagccagg 720 780 acggccattg gccacgaccg gaaagggcag ctggtgctct ttcatgcaga cggccatacg gagcagcgtg gcatcaacct gtgggaaatg gcggagttcc tgctgaaaca ggacgtggtc 840 900 aacgccatca acctggatgg gggtggctct gccacctttg tgctcaacgg gaccttggcc agttaccegt cagatcactg ccaggacaac atgtggcgct gtccccgcca agtgtccacc 960 gtggtgtgtg tgcacgaacc ccgctgccag ccgcctgact gccacggcca cgggacctgc 1020 gtggacgggc actgccaatg caccgggcac ttctggcggg gtcccggctg tgatgagctg 1080

| gactgtggcc | cctctaactg | cagccagcac | ggactgtgca | cggagaccgg | ctgccgctgt | 1140 |
|------------|------------|------------|------------|------------|------------|------|
| gatgccggat | ggaccgggtc | caactgcagt | gaagagtgtc | cccttggctg | gcatgggccg | 1200 |
| ggctgccaga | ggcgttgtaa | gtgtgagcac | cattgtccct | gtgaccccaa | gactggcaac | 1260 |
| tgcagcgtct | ccagagtaaa | gcagtgtctc | cagccacctg | aagccaccct | gagggcggga | 1320 |
| gaactctcct | ttttcaccag | gaccgcctgg | ctagccctca | ccctggcgct | ggccttcctc | 1380 |
| ctgctgatca | gcattgcagc | aaacctgtcc | ttgctcctgt | ccagagcaga | gaggaaccgg | 1440 |
| cgcctgcatg | gggactatgc | ataccacccg | ctgcaggaga | tgaacgggga | gcctctggcc | 1500 |
| gcagagaagg | agcagccagg | gggcgcccac | aaccccttca | aggactgaag | cctcaagctg | 1560 |
| cccggggtgg | cacgtcgcga | aagcttgttt | ccccacggtc | tggcttctgc | aggggaaatt | 1620 |
| tcaaggccac | tggcgtggac | catctgggtg | tcctcaatgg | cccctgtggg | gcagccaagt | 1680 |
| tcctgatagc | acttgtgcct | cageceetca | cctggccacc | tgccagggca | cctgcaaccc | 1740 |
| tagcaatacc | atgctcgctg | gagaggctca | gctgcctgct | tctcgcctgc | ctgtgtctgc | 1800 |
| tgccgagaag | cccgtgcccc | cgggagggct | gccgcactgc | caaagagtct | ccctcctcct | 1860 |
| ggggaagggg | ctgccaacga | accagactca | gtgaccacgt | catgacagaa | cagcacatcc | 1920 |
| tggccagcac | ccctggctgg | agtgggttaa | agggacgagt | ctgccttcct | ggctgtgaca | 1980 |
| cgggacccct | tttctacaga | cctcatcact | ggatttgcca | actagaattc | gatttcctgt | 2040 |
| cataggaagc | tccttggaag | aagggatggg | gggatgaaat | catgtttaca | gacctgtttt | 2100 |
| gtcatcctgc | tgccaagaag | ttttttaatc | acttgaataa | attgatataa | taaaaggagc | 2160 |
| caccaggtgg | tgtgtggatt | ctg | | | | 2183 |

<210> 18

Met Ala Thr Ser Thr Gly Arg Trp Leu Leu Leu Arg Leu Ala Leu Phe 1 5 10 15

Gly Phe Leu Trp Glu Ala Ser.Gly Gly Leu Asp Ser Gly Ala Ser Arg 20 25 30

Asp Asp Leu Leu Leu Pro Tyr Pro Arg Ala Arg Ala Arg Leu Pro 35 40 45

<211> 515

<212> PRT

<213> Homo sapiens

<400> 18

Arg Asp Cys Thr Arg Val Arg Ala Gly Asn Arg Glu His Glu Ser Trp 50 55 60

Pro Pro Pro Pro Ala Thr Pro Gly Ala Gly Gly Leu Ala Val Arg Thr 65 70 75 80

Phe Val Ser His Phe Arg Asp Arg Ala Val Ala Gly His Leu Thr Arg 85 90 95

Ala Val Glu Pro Leu Arg Thr Phe Ser Val Leu Glu Pro Gly Gly Pro 100 105 110

Gly Gly Cys Ala Ala Arg Arg Arg Ala Thr Val Glu Glu Thr Ala Arg 115 120 125

Ala Ala Asp Cys Arg Val Ala Gln Asn Gly Gly Phe Phe Arg Met Asn 130 135 140

Ser Gly Glu Cys Leu Gly Asn Val Val Ser Asp Glu Arg Arg Val Ser 145 150 155 160

Ser Ser Gly Gly Leu Gln Asn Ala Gln Phe Gly Ile Arg Arg Asp Gly . 165 170 175

Thr Leu Val Thr Gly Tyr Leu Ser Glu Glu Glu Val Leu Asp Thr Glu 180 185 190

Asn Pro Phe Val Gln Leu Leu Ser Gly Val Val Trp Leu Ile Arg Asn 195 200 205

Gly Ser Ile Tyr Ile Asn Glu Ser Gln Ala Thr Glu Cys Asp Glu Thr 210 215 220

Gln Glu Thr Gly Ser Phe Ser Lys Phe Val Asn Val Ile Ser Ala Arg 225 230 235 240

Thr Ala Ile Gly His Asp Arg Lys Gly Gln Leu Val Leu Phe His Ala 245 250 255

Asp Gly His Thr Glu Gln Arg Gly Ile Asn Leu Trp Glu Met Ala Glu 260 265 270

Phe Leu Lys Gln Asp Val Val Asn Ala Ile Asn Leu Asp Gly Gly 275 280 285

Gly Ser Ala Thr Phe Val Leu Asn Gly Thr Leu Ala Ser Tyr Pro Ser 290 295 300

Asp His Cys Gln Asp Asn Met Trp Arg Cys Pro Arg Gln Val Ser Thr 305 310 315 320

Val Val Cys Val His Glu Pro Arg Cys Gln Pro Pro Asp Cys His Gly 325 330 335

His Gly Thr Cys Val Asp Gly His Cys Gln Cys Thr Gly His Phe Trp 340 345 350

Arg Gly Pro Gly Cys Asp Glu Leu Asp Cys Gly Pro Ser Asn Cys Ser 355 360 365

Gln His Gly Leu Cys Thr Glu Thr Gly Cys Arg Cys Asp Ala Gly Trp 370 375 380

Thr Gly Ser Asn Cys Ser Glu Glu Cys Pro Leu Gly Trp His Gly Pro 385 390 395 400

Gly Cys Gln Arg Arg Cys Lys Cys Glu His His Cys Pro Cys Asp Pro 405 410 415

Lys Thr Gly Asn Cys Ser Val Ser Arg Val Lys Gln Cys Leu Gln Pro 420 425 430

Pro Glu Ala Thr Leu Arg Ala Gly Glu Leu Ser Phe Phe Thr Arg Thr 435 440 445

Ala Trp Leu Ala Leu Thr Leu Ala Leu Ala Phe Leu Leu Leu Ile Ser 450 460

Ile Ala Ala Asn Leu Ser Leu Leu Leu Ser Arg Ala Glu Arg Asn Arg 465 470 475 480

Arg Leu His Gly Asp Tyr Ala Tyr His Pro Leu Gln Glu Met Asn Gly
485 490 495

Glu Pro Leu Ala Ala Glu Lys Glu Gln Pro Gly Gly Ala His Asn Pro 500 505 510

Phe Lys Asp 515

<210> 19

<211> 2005

<212> DNA

<213> Mus musculus

<400> 19 60 gtttcccgcg acgatgacct gctgctgcct tacccactag cgcgcagacg tccctcgcga gactgcgccc gggtgcgctc aggtagccca gagcaggaga gctggcctcc gccacctctg 120 gccacccacg aaccccgggc gccaagccac cacgcggccg tgcgcacctt cgtgtcgcac 180 ttcgagggc gcgcggtggc cggccacctg acgcgggtcg ccgatcccct acgcactttc 240 tcggtgctgg agcccggagg agccgggggc tgcggcggca gaagcgccgc ggctactgtg 300 gaggacacag ccgtccgggc cggttgccgc atcgctcaga acggtggctt cttccgcatg 360 agcactggcg agtgcttggg gaacgtggtg agcgacgggc ggctggtgag cagctcaggg 420 480 ggactgcaga acgcgcagtt cggtatccga cgcgatggaa ccatagtcac cgggtcctgt cttgaagaag aggttetgga teeegtgaat eegttegtge agetgetgag eggagtegtg 540 tggctcatcc gcaatggaaa catctacatc aacgagagcc aagccatcga gtgtgacgag 600 acacaggaga caggttettt tagcaaattt gtgaatgtga tgteageeag gacageegtg 660 ggtcatgacc gtgaggggca gcttatcctc ttccatgctg atggacagac ggaacagcgt 720 ggccttaacc tatgggagat ggcagagttc ctgcgtcaac aagatgtcgt caatgccatc 780 aacctggatg gaggcggttc tgctactttt gtgctcaatg ggaccctggc cagttaccct 840 tcagatcact gccaggacaa catgtggcgc tgtccccgcc aagtgtccac tgtggtgtgt 900 gtgcatgaac cgcgctgcca gccacccgac tgcagtggcc atgggacctg tgtggatggc 960 cactgtgaat gcaccagcca cttctggcgg ggcgaggcct gcagcgagct ggactgtggc 1020 ccctccaact gcagccagca tgggctgtgc acagctggct gccactgtga tgctgggtgg 1080 1140 acaggatcca actgcagtga agagtgtcct ctgggctggt atgggccagg ttgccagagg ccctgccagt gtgagcacca gtgtttctgt gacccgcaga ctggcaactg cagcatctcc 1200 caagtgaggc agtgtctcca gccaactgag gctacgccga gggcaggaga gctggcctct 1260 ttcaccagga ccacctggct agccctcacc ctgacactaa ttttcctgct gctgatcagc 1320

| actggggtca | acgtgtcctt | gttcctgggc | tccagggccg | agaggaaccg | gcacctcgac | 1380 |
|------------|------------|------------|------------|------------|------------|------|
| ggggactatg | tgtatcaccc | actgcaggag | gtgaacgggg | aagcgctgac | tgcagagaag | 1440 |
| gagcacatgg | aggaaactag | caaccccttc | aaggactgaa | gagctgcccc | aacggcatgc | 1500 |
| tccagataat | cttgtccctg | ctcctcactt | ccacagggga | cattgtgagg | ccactggcat | 1560 |
| ggatgctatg | caccccaccc | tttgctggcc | atattcctcc | tgtccccatg | ctgtggctca | 1620 |
| tgccaaccta | gcaataagga | gctctggaga | gcctgcacct | gcctcccgct | cgcctatatc | 1680 |
| tgctgcccag | aggcctgtct | cgcacagggg | tctcgccact | gccaaagact | cccaggaagt | 1740 |
| caaagactcc | cagtaatcca | ctagcaaatg | gaactctgta | acgccatcat | aacaagagtg | 1800 |
| gccactctcc | gcgtgcacag | gtatgaaata | taaatcctta | cacacacaca | cacacacacc | 1860 |
| ctcggctcag | ccacggcact | cgccttttat | acagcgtcat | cgctggacag | ccaactagaa | 1920 |
| ctctgcatcc | tgtcacagga | agcacctcat | aagaaggaat | ggggagggaa | ggcagtcgcc | 1980 |
| ttgttttcag | accttagccg | aattc | | | | 2005 |

<210> 20 <211> 492 <212> PRT <213> Mus musculus

<400> 20

Val Ser Arg Asp Asp Asp Leu Leu Leu Pro Tyr Pro Leu Ala Arg Arg 1 5 10 15

Arg Pro Ser Arg Asp Cys Ala Arg Val Arg Ser Gly Ser Pro Glu Gln 20 25 30

Glu Ser Trp Pro Pro Pro Pro Leu Ala Thr His Glu Pro Arg Ala Pro 35 40 45

Ser His His Ala Ala Val Arg Thr Phe Val Ser His Phe Glu Gly Arg 50 55 60

Ala Val Ala Gly His Leu Thr Arg Val Ala Asp Pro Leu Arg Thr Phe 65 70 75 80

Ser Val Leu Glu Pro Gly Gly Ala Gly Gly Cys Gly Gly Arg Ser Ala 85 90 95

Ala Ala Thr Val Glu Asp Thr Ala Val Arg Ala Gly Cys Arg Ile Ala 100 105 110

Gln Asn Gly Gly Phe Phe Arg Met Ser Thr Gly Glu Cys Leu Gly Asn 115 120 125

Val Val Ser Asp Gly Arg Leu Val Ser Ser Ser Gly Gly Leu Gln Asn 130 135 140

Ala Gln Phe Gly Ile Arg Arg Asp Gly Thr Ile Val Thr Gly Ser Cys 145 150 155 160

Leu Glu Glu Val Leu Asp Pro Val Asn Pro Phe Val Gln Leu Leu 165 170 175

Ser Gly Val Val Trp Leu Ile Arg Asn Gly Asn Ile Tyr Ile Asn Glu 180 185 190

Ser Gln Ala Ile Glu Cys Asp Glu Thr Gln Glu Thr Gly Ser Phe Ser 195 200 205

Lys Phe Val Asn Val Met Ser Ala Arg Thr Ala Val Gly His Asp Arg 210 215 220

Glu Gly Gln Leu Ile Leu Phe His Ala Asp Gly Gln Thr Glu Gln Arg 225 230 235 240

Gly Leu Asn Leu Trp Glu Met Ala Glu Phe Leu Arg Gln Gln Asp Val 245 250 255

Val Asn Ala Ile Asn Leu Asp Gly Gly Gly Ser Ala Thr Phe Val Leu 260 265 270

Asn Gly Thr Leu Ala Ser Tyr Pro Ser Asp His Cys Gln Asp Asn Met 275 280 285

Trp Arg Cys Pro Arg Gln Val Ser Thr Val Val Cys Val His Glu Pro 290 295 300

Arg Cys Gln Pro Pro Asp Cys Ser Gly His Gly Thr Cys Val Asp Gly 305 310 315 320

His Cys Glu Cys Thr Ser His Phe Trp Arg Gly Glu Ala Cys Ser Glu

| I | .eu | Asp | Cys | Gly 340 | Pro | Ser | Asn | Cys | Ser 345 | Gln | His | Gly | Leu | Cys 350 | Thr | Ala | | |
|---|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|---|-----|
| c | Sly | Cys | His 355 | Cys | Asp | Ala | Gly | Trp 360 | Thr | Gly | Ser | Asn | Cys 365 | Ser | Glu | Glu | | |
| C | Cys | Pro 370 | Leu | Gly | Trp | Tyr | Gly 375 | Pro | Gly | Cys | Gln | Arg 380 | Pro | Cys | Gln | Сув | | |
| | Glu 885 | His | Gln | Cys | Phe | Cys 390 | Asp. | Pro | Gln | Thr | Gly 395 | Asn | Cys | Ser | Ile | Ser 400 | | |
| C | Gln | Val | Arg | Gln | Cys 405 | Leu | Gln | Pro | Thr | Glu 410 | Ala | Thr | Pro | Arg | Ala 415 | Gly | | |
| C | Glu | Leu | Ala | Ser 420 | Phe | Thr | Arg | Thr | Thr 425 | Trp | Leu | Ala | Leu | Thr 430 | Leu | Thr | | |
|] | Leu | Ile | Phe 435 | | Leu | Leu | Ile | Ser 440 | | Gly | Val | Asn | Val 445 | Ser | Leu | Phe | | |
|] | Leu | Gly 450 | | Arg | Ala | Glu | Arg 455 | | Arg | His | Leu | Asp 460 | | Asp | Tyr | Val | | |
| | Tyr 465 | | Pro | Leu | Gln | Glu 470 | | · Asn | Gly | Glu | Ala 475 | | Thr | Ala | Glu | Lys 480 | | |
| , | Glu | His | Met | Glu | Glu 485 | | Ser | Asn | Pro | Phe 490 | - | Asp | , | | | | | |
| | | 1> 2> | | e musc | ulus | ı | | | | | | | | | | | | |
| | <40 cag | | 21 :999 | actt | acta | ıta a | caca | ıggac | a ct | tgto | acct | : gaa | agct | tga | gtca | gtcagt | | 60 |
| | | | | | | | | | | | | | | | | gtagat | | 120 |
| | cag | gactt | tct | acag | gccaa | itt c | tctt | ctto | c to | ctct | ccat | . ggg | gttca | aggg | tctt | catctc | : | 180 |

aggttgcaca gcgagttcat ttatgtgctg tgccatctcg ccagtcgttc ctatatccta

300 gaggaaaact agtttcttct ggtcaagagg aggaaagagt ggagacctgt cattctaaga 360 tacccaaaac agggccaggt tggggacctg tgcctttaat cccatcactt ggggattagg 420 tagaagcaag aggetetaga eeagtetaca eactgaattt eaageeagee tacetataaa tcagagaccc tgcttcaaaa ataaaattaa acaaaaacga agataaacca agctacccaa 480 aacacaagag ttaatccagt cagacaggtc tagcaaatgc taggatgaaa ggtgtgcacc 540 accacgagtg ggctgcaagc ctctctctct ctctctctct ctctctctct ctcgtttgtt 600 ttgttttteg agacaaggtt tetetgtgta gecetggetg teetggaact caetetgtag 660 accaggetgg cetegagett caetettaaa agtteetett ceteeteete catettttee 720 tectettace ecctaggete etttteetet tettgtettt eagataaagt eteaagtagt 780 ccagactggt ctcaaactaa ctaactagcc aagaatagcc aacctcttaa cttccgattc 840 900 teetgeetet getgaatget ggggttgtgg egtgggeeae eaettetggt ttgtgeaaea cagaaggaac tagggcttta agcacgagaa gcaagttctg tacagactta cacaggccca 960 1020 gcatctgttc ttgcaatttt ctgtaagttt gacataatat gagaataaaa agctatctat ctcccttcca gccttaccct ctctgatgga attcgaatgc gtaatcaaag cacccaacag 1080 cctggcctga aatcacgtgg ggcaagccca cgtgaccgga gcaccaatcc aatatggcgg 1140 1200 egeceagggg gecegggetg tteeteatac eegegetget eggettacte ggggtggegt 1260 ggtgcagctt aagcttcggg tgagtgcaag ccgccggggc cagcctggct ggggtccacc ttteetgage geteteagge acagecetee gaceteaega tegeceegte eetgeagggt 1320 1380 ttcccgcgac gatgacctgc tgctgcctta cccactagcg cgcagacgtc cctcgcgaga ctgcgcccgg gtgcgctcag gtagcccaga gcaggagagc tggcctccgc cacctctggc 1440 cacccacgaa ccccgggcgc caagccacca cgcggccgtg cgcaccttcg tgtcgcactt 1500 egaggggege geggtggeeg gecaeetgae gegggtegee gateeeetae geaetttete 1560 1620 ggtgctggag cccggaggag ccgggggctg cggcggcaga agcgccgcgg ctactgtgga ggacacagcc gtccgggccg gttgccgcat cgctcagaac ggtggcttct tccgcatgag 1680 cactggcgag tgcttgggga acgtggtgag cgacgggcgg ctggtgagca gctcaggggg 1740 actgcagaac gcgcagttcg gtatccgacg cgatggaacc atagtcaccg ggtgaggagg 1800 cagggagccc cggggctgta gagggcaaag ggtctctgat gttctttcag agccatgcct 1860 ecgagtecag gtecetaace aaactteetg tetttettet teegagtaat gaegetgaea 1920 cetteettee tttaagttta tteatgtgee actgaataat etgtgateag geegtgtgtg 1980

gggacttggg gaggcgaccg tgagcctgaa cacagtttgt gccctagtga actttgtgta 2040 2100 gtattagaga aacatttcgt gttcaacgaa gccatggaac caattggaaa tagtgtagag 2160 tttatggagc agtcccagac agctagctgg aggccttttg ctgtcctgat aaaaatccag 2220 gttagacaag gagcttgttg agggcagcct ttggaagttt ctgtgtttct tgaaatttga cagcagccag agttgacagc aggcaggcag gagtagaagg tagcgccatc tggtgttcca 2280 2340 gttctcttcc aaggttccgt tttttgccaa ggctgggaag tgggctttcc ccaactcttc tragccettg gttgcaattt ctgggcetge ceatgtatet ggttetteat cetteaacat 2400 2460 cagccagtgt caccactgtt gatcttaggt tttcacagat cctaaaactt ctgccagtga ccagegeetg cagtttetet tecetggete tgteetteaa eetetetaca ttecageeat 2520 ctccctagct cctctcttgg actccctttc agacttgttg tcatgatcac tgtctcagaa 2580 2640 cccctattgc tcctttacaa tggtccactg acctgctcac ctcctacttt tttttttaa 2700 atgtgtgtgc atctgtgtgt gcctgagggg agaccagagt ttgatttcaa atgtcttcta ttototttto otocatotta ttttotaaca caaaatotga atotagagat cactggttoa 2760 2820 2880 cagcactaag getacagtgt gtgctgttcc agccagcttt ctcatgggtg ctgaggatct 2940 gaacgcaggt tcacatgtgt ggtgggaagg cttttaccca atgctctgtc tttccagccc atcctccctt gttaactgcc aaacagctgc ctatcctgtc catgtgtagc tcactgctac 3000 3060 ttcttttatt atgaggtcag cacatgttac taaagatggc aagagaagaa ggttctttca 3120 ttgtgtcata gctatagctc aggaggaatt ttatttcctg tgtaggcaca caggagagca tettecaget cacactecaa etgaactaae tgaacacetg eetatatate caaagaaggg 3180 gtgtcagtgc caatcacagc acacctccag tgcaaatgaa ggtttgtgtt tgcaccaatc 3240 3300 acageettge etetttage atgeateaca acaaagteet eetagaetat eaggggatat 3360 gctctcttgg ccaaggtagg aatagttgca gtgtcatctg gcacaaacca tttcaaacgg cctggctgag gttatgcctt cgggaacctg aagtetttgt gtggttgtet ccaagtgtet 3420 3480 gtggagetee aggeggetgg tgetgaeaga egetttgtet agttggetgt ttgaettttg cttaagcagc cagggcagta gagtctaaca gatgctaatt tcaggatcag gaagactgta 3540 3600 gaaaaatgag catcaagaag cccctggtac ccaaagctgc tcttgccaat gagtgaacct ctgccttccc gcttccaggt cctgtcttga agaagaggtt ctggatcccg tgaatccgtt 3660

3720 cgtgcagctg ctgagcggag tcgtgtggct catccgcaat ggaaacatct acatcaacga gagccaagcc atcgagtgtg acgagacaca ggagacaggt caggaagcac aggtgttctg 3780 ttttatttgt attaggtttt gatttgttta ttttgtgcat gcagcgggtg catgcatgct 3840 cctttccttt cgccatgtga gtcctgagta ttgaactcag actgttaagt gtgatgggag 3900 3960 gcactttacc cactgagcca ctttcccagc cctcagcatc agctttcttc agacccagga 4020 acagtgtgag tgggttattc tttagtgttc ccaaacattt actgagcagc tatttactgt ttagcactat ggtgagagtc ctagggattc agtcttatgt agaatataga aggagaatcc 4080 ttggcaataa gctggaaaat tgtgacaagt gccaagaaag aaacaggaga aaggggaccg 4140 gtggggacca gaagcacagg tatgaggaaa gtgcctgcag atttgctgta tggtggcctc 4200 cacatggcct aggagtttgt cataaatgca gagccatgag tccaccctcc ctatacctcc 4260 4320 catccagaaa ccactggtta aatcctaaca acttgggtgt gcaggcactc ccttggtgac tetgatggae acteaaggte aagggeeact tggggatggg etgatgagtt ggettggtea 4380 4440 gtaaagtatt tgccttgaaa gtgtgaggac ctgagttgga gccccagaaa gaaacattaa 4500 aagccaagtg ctgggatgca cacttgcatt cccagggatg gagctggaag gcagggatag gcagatccac ggccacacgg tgatattcta agctaacaag agacctgtct cacacagaaa 4560 gtgggtggca cctgaggacc aacacccagg gttatcctct gacgtacctc cagagtggaa 4620 aatactgggg tggtggaaaa ggacactttg gtcctgggaa tctggctatt cagggtatag 4680 tgtagaggga gagggagact caagaggctg tctttgagtc aaaggaacaa gctatcagaa 4740 gaactcaggg cagaggcctg tggttcccag gctcagggca gccttcaagg ccctaggcag 4800 4860 agagtagctg ctgggtgaac aagtacagaa gtgaggcctg gggcctcagg caaggcctgt gaaatccttc caccaacata gaagtttctg gagactgaga tcacatgaag tgcttctggc 4920 4980 tgtggcatgg aagctcactg gaggtggagc tgggatgtgg ctcagtgatc cagtgcttgc cacacgtgca cgagggaagg agccatcaaa agagagaaag tcgggagacc tgaggggtcc 5040 5100 cctggagagc tgggtaacca ccccgggccc ttctccttta ggttctttta gcaaatttgt 5160 gaatgtgatg tcagccagga cagccgtggg tcatgaccgt gaggggcagc ttatcctctt ccatgctgat ggacagacgg aacagcgtgg tgagtcccag gaaccttggg gctgtttgca 5220 cttcagccac cctacctttc cagtcggttc tggggtattg gtgggacaag acagctttcc 5280 ggccattttg gaagtttcat ctggaggcaa tagcatttac ctactagtga aagaagccag 5340 ttaagccaga gaccacaggg gctcaagctg cataccccct ctgcacagcc ttaacctatg 5400 ggagatggca gagttcctgc gtcaacaaga tgtcgtcaat gccatcaacc tggatggagg 5460 cggttctgct acttttgtgc tcaatgggac cctggccagt tacccttcag atcactggta 5520 agaaccettg agecacettt gtggetetet cagaetgtet caeteagtea ataetgagae 5580 5640 cctgttgtgt gccaggccct gggtatccaa aagtgagcag aagagccgag atctcttccc tcagggtgct gcacagccca tccctggaaa cctgagacag gtcaggaaag gcctccctga 5700 ggacagtgaa gtaagacctg aggagatggc tggccggggt tgagagagcc tttaccggaa 5760 gacaaactgt acgcaatggg gaaatccgct aagtggccca gggagaggct ggagctatag 5820 ctcaggagga aaagtacttg cctcgcaagc gaaggacctg agtttaaact ccaaaaccca 5880 tataaaaagc cagatacgag caagtggcac atgcttgcag teccageett gttgaggaag 5940 agtcaggtga atcctgaccc tctggccagc cagcctagcc tactttttgg caaggtccag 6000 gccagcgaga aagataaata aaataaagtt ttaaatgaca tgtatctaag gttgtcctga 6060 6120 ctccatatgc gcacgcacgc atgcacgcac gcacaactgg cagaatggaa agggaggcaa actggacage ctttatagge tgeggeaggg accageacea aggeetagae etegteteae 6180 agtgaatccc ccacagecag gacaacatgt ggcgctgtcc ccgccaagtg tccactgtgg 6240 6300 tgtgtgtgca tgaaccgcgc tgccagccac ccgactgcag tggccatggg acctgtgtgg atggccactg tgaatgcacc agccacttct ggcggggcga ggcctgcagc gagctggact 6360 gtggcccctc caactgcagc cagcatgggc tgtgcacaga gagtgagtgg ggagcccaca 6420 ggagggtggt gctctggcgg gaccccagct cgcccatgct agactcccgc ctgtgtcctt 6480 acccagcete tgtggtettg etttggtage tggetgeeae tgtgatgetg ggtggaeagg 6540 6600 atccaactgc agtgaaggtg agagctgcct gcaaacactc ctggagaggg tggcctggct gcacgcagct ggtatgacgc cttcgtccct ccttctggct tggaacttac cttcagagcc 6660 ttttctcatt tcgcatgtgg atacccgatg ttctacctac tgaaagagcc cacaagtagg 6720 6780 aagccagatt ttcagtattg tcactcaact ctaaggacca atagcaaaaa aacaaagtgg 6840 ccacgccct gagggagatc caccaaagtc cttaactcct ggaaagcagc tcctggtgat cctaggcatg ggtagggtgg tttcagcatc agctcagtgg agttcccatt cataatttct 6900 tcatcctttt aaggtcataa gttctagagc ccaccttaaa tctaggcagt attcttggtg 6960 tttatctgag acaaagtctt atacagccca cgcagttctc taacttagta tgtaaccgag 7020 aatggcctca agcaacctgc ttcctccttt caagcgctgg gattataggc atagcaccaa 7080



gccacctcac ccagccctgc tggacttaaa ttgggtcttc attttataag acaagcatga 8880 gctaattccc cagttcctaa aatgttttta acatccttaa acatcagaga ctgtctgtgg 8940 9000 cccagtttgg gttctggctg cctgccccca ctcaagactc tcttttccat ctcaggacca 9060 cctggctagc cctcaccctg acactaattt tcctgctgct gatcagcact ggggtcaacg 9120 9180 tgtccttgtt cctgggctcc agggccgaga ggaaccggca cctcgacggg gactatgtgt 9240 atcacccact gcaggaggtg aacggggaag cgctgactgc agagaaggag cacatggagg aaactagcaa ccccttcaag gactgaagag ctgccccaac ggcatgctcc agataatctt 9300 gtccctgctc ctcacttcca caggggacat tgtgaggcca ctggcatgga tgctatgcac 9360 9420 cccaccettt getggecata tteeteetgt ceceatgetg tggeteatge caacctagea 9480 ataaggaget etggagagee tgeacetgee teeegetege etatatetge tgeecagagg 9540 cctqtctcqc acaqqqqtct cqccactqcc aaagactccc aggaagtcaa agactcccag taatccacta gcaaatggaa ctctgtaacg ccatcataac aagagtggcc actctccgcg 9600 tgcacaggta tgaaatataa atccttacac acacacaca acacaccctc ggctcagcca 9660 eggeactege ettttataca gegteatege tggacageca actagaacte tgcateetgt 9720 9780 cacaggaagc acctcataag aaggaatggg gagggaaggc agtcgccttg ttttcagacc 9792 ttagccgaat tc

```
<210>
       22
<211>
<212>
       PRT
<213>
       Artificial Sequence
<220>
<223>
      synthetic peptide
<400>
       22
Ile Glu Gly Arg
<210>
       23
<211>
<212>
       PRT
<213>
       Artificial Sequence
<220>
<223> synthetic peptide
```

```
<400> 23
Asp Asp Asp Lys
<210> 24
<211> 6
<212> PRT
<213> Artificial Sequence
<223> synthetic peptide
<400> 24
Arg Ala Arg Tyr Lys Arg
            5
<210> 25
<211> 6
<212> PRT
<213> Artificial Sequence
<220>
<223> synthetic peptide
<400> 25
Pro Gly Ala Ala His Tyr
      5
<210> 26
<211> 46
<212> PRT
<213> Homo sapiens
<400> 26
Asp Glu Glu Ser Leu Lys Thr Gln Leu Ala Tyr Phe Thr Asp Ser
Lys Asn Thr Gly Arg Gln Leu Lys Asp Thr Phe Ala Asp Ser Leu Arg
Tyr Val Asn Lys Ile Leu Asn Ser Lys Phe Gly Phe Thr Ser
<210> 27
<211> 112
<212> DNA
```

| <213> | Homo sapiens | |
|--------|--|-----|
| <400> | 27 | |
| gacgaa | agaag agtcattgaa gacacaattg gcatacttca ctgatagcaa gaatactggg | 60 |
| aggcaa | actaa aagatacatt tgcagattcc ctcagatatg taaataaaat tc | 112 |
| <210> | 28 | |
| <211> | | |
| <212> | | |
| <213> | Artificial Sequence | |
| <220> | | |
| <223> | synthetic DNA | |
| <400> | 28 | |
| ctcagt | aact tetgtgttaa eeggatgaae eggaattaat tettatgae | 49 |
| <210> | 29 | |
| <211> | 48 | |
| <212> | DNA | |
| <213> | Artificial Sequence | |
| <220> | | |
| <223> | synthetic DNA | |
| <400> | 29 | |
| gaggca | atta attaacgtet etgeagatte eetcagatat gtaaataa | 48 |
| | | |
| <210> | 30 | |
| <211> | 11 | |
| <212> | PRT | |
| <213> | Homo sapiens | |
| <400> | 30 | |
| Asp Gl | u Glu Glu Ser Leu Lys Thr Gln Leu Ala | |
| 1 | 5 10 | |
| | | |
| <210> | 31 | |
| <211> | 10 | |
| <212> | PRT | |
| <213> | hybrid | |
| <400> | 31 | |
| | p Ser Leu Arg Tyr Val Asn Lys Ile | |
| 1 | 5 10 | |
| <210> | 32 | |
| <211> | 88 | |
| <212> | DNA | |
| | | |

| <213> hybrid | |
|---|----|
| <400> 32 gacgaagaag agtcattgaa gacacaattg gcctacttgg ccttaattaa cgtctctgca | 60 |
| gattccctca gatatgtaaa taaaattc | 88 |
| <210> 33 <211> 25 <212> PRT <213> hybrid | |
| <400> 33 | |
| Thr Gln Leu Ala Tyr Phe Thr Asp Ser Lys Asn Thr Gly Ile Glu Gly 1 5 10 15 | |
| Arg Asp Thr Phe Ala Asp Ser Leu Arg 20 25 | |
| <210> 34 <211> 75 <212> DNA <213> hybrid | |
| <400> 34 acacaattgg cctacttcac tgatagcaag aatactggga tcgagggaag agatacattt | 60 |
| ggagattagg tagg | |
| gcagattece teaga | 75 |
| <210> 35 <211> 25 <212> PRT <213> hybrid | 75 |
| <210> 35 <211> 25 <212> PRT | 75 |
| <210> 35 <211> 25 <212> PRT <213> hybrid | 75 |
| <pre> <210> 35 <211> 25 <212> PRT <213> hybrid <400> 35 Thr Gln Leu Ala Tyr Phe Thr Asp Ser Lys Asn Arg Ala Arg Tyr Lys</pre> | 75 |
| <pre> <210> 35 <211> 25 <212> PRT <213> hybrid <400> 35 Thr Gln Leu Ala Tyr Phe Thr Asp Ser Lys Asn Arg Ala Arg Tyr Lys 1</pre> | 75 |
| <pre> <210> 35 <211> 25 <212> PRT <213> hybrid <400> 35 Thr Gln Leu Ala Tyr Phe Thr Asp Ser Lys Asn Arg Ala Arg Tyr Lys 1</pre> | 75 |

acacaattgg cctacttcac tgatagcaag aatagagcca gatacaagag agatacattt 60

gcagattccc tcaga

75

<210> 37 <211> 25 <212> PRT <213> hybrid <400> 37

Thr Gln Leu Ala Tyr Phe Thr Asp Ser Lys Asn Thr Asp Asp Asp Asp 1 5 10 15

Lys Asp Thr Phe Ala Asp Ser Leu Arg
20 25

<210> 38 <211> 25 <212> PRT <213> hybrid <400> 38

Thr Gln Leu Ala Tyr Phe Thr Asp Ser Lys Asn Pro Gly Ala Ala His 1 5 10 15

Tyr Asp Thr Phe Ala Asp Ser Leu Arg 20 25